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Article

# Examining the Relationships between the Incidence of Infectious Diseases and Mood Disorders: An Analysis of Data from the Global Burden of Disease Studies, 1990–2019

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Abstract: Mood disorders are among the commonest mental disorders worldwide. Epidemiological and clinical evidence suggests that there are close links between infectious diseases and mood disorders, but the strength and direction of these association remain largely unknown. Theoretical models have attempted to explain this link based on evolutionary or immune-related factors, but these have not been empirically verified. The current study examined cross-sectional and longitudinal associations between the incidence of infectious diseases and mood disorders, while correcting for climate and economic factors, based on data from the Global Burden of Disease Studies, 1990-2019. It was found that major depressive disorder was positively associated with lower respiratory infections, while bipolar disorder was positively associated with upper respiratory infections and negatively associated with enteric and tropical infections, both cross-sectionally and over a period of thirty years. These results suggest that a complex, bidirectional relationship exists between these disorders. This relationship may be mediated through the immune system as well as through the gut-brain and lung-brain axes. Understanding the mechanisms that link these groups of disorders could lead to advances in the prevention and treatment of both.

**Keywords:** major depressive disorder; bipolar disorder; infectious diseases; upper respiratory infections; lower respiratory infectious; lung-brain axis; gut-brain axis

# 1. Introduction

Major depressive disorder (MDD), commonly referred to as depression, is the most common mental disorder globally. According to an estimate published by the World Health Organization in 2018, the point prevalence of MDD is around 4.4% of the world's population; in other words, over 300 million people around the world suffer from depression at any given time [1]. Data from the World Mental Health Survey Initiative yielded somewhat higher estimates, with a 12-month prevalence of around 5-6% and a lifetime prevalence of 11-14% [2]. Due to its frequency, symptomatology, and tendency to recur or persist, depression is one of the leading global causes of disability [3]. A significant proportion of patients with an initial diagnosis of MDD may later be rediagnosed as having bipolar disorder (BD). BD is characterized by recurrent episodes of both elevated mood (mania) and depression, and is associated with a higher level of disability than MDD [4]. A recent meta-analysis has found that over 22% of patients with a diagnosis of MDD subsequently develop BD, with the risk of this "transition" being highest in the first five years after an episode of MDD [5]. Though BD and MDD are considered to be distinct disorders in current classificatory systems, there is a substantial clinical, genetic and neurobiological overlap between them. These disorders are probably best understood as different manifestations of an underlying genetic vulnerability towards mood disorder [6-8]. A more fine-grained analysis suggests that depression may overlap genetically with bipolar type II disorder (BD-II), a condition in which episodes of



elevated mood are milder ("hypomania"), while more severe episodes of elevated mood ("mania") may represent a distinct genetic diathesis [9].

There are significant cross-national variations in the epidemiology of MDD, with higher prevalence estimates being reported in high-income countries than in low- and middle-income countries [2,10]. This finding appears paradoxical, as it is expected that the stressful life events that trigger depression or contribute to its persistence would be more common in the context of social unrest and economic hardship [11,12]. Several explanations have been advanced to account for this result, including differences in population genetics [13,14], cultural values and practices [15], diet [16,17], levels of urbanization [18], income inequality [19], and even specific attitudes or stressors seen in higher-income groups [20]. Some researchers have even considered depression a "disease of civilization" linked to several aspects of the "modern" lifestyle. These explanations are not mutually exclusive, and several potential links between them have been identified. Some researchers have highlighted a process of "co-evolution" between population genetics and cultural values which correlates with cross-national variations in the prevalence of MDD [21,22], while others have documented evidence of interactions between specific vulnerability genes, such as *IL6*, and exposure to social stress as contributing to variations in depressive symptoms across the lifespan [23,24].

Among the risk factors that contribute to variations in the risk of depression, the link between infectious diseases and MDD has attracted increasing attention in the past two decades. From an evolutionary perspective, similarities have been noted between the symptoms of depression and "sickness behavior", which has evolved as a defence against infectious diseases. Sickness behavior, which is mediated by cytokines such as interleukin-6 (IL-6) and tumor necrosis factor alpha (TNF- $\alpha$ ) includes reduced appetite, fatigue, reduced physical activity and social withdrawal. This set of responses is adaptive both at the individual level (by facilitating tissue healing and recovery from infectious diseases) and at the group level (by minimizing transmission to other members of the same species) [25–28]. These similarities, as well as the fact that MDD is associated with elevated levels of these cytokines [29], have prompted several researchers to investigate the possibility of an immune-mediated link between exposure to infectious diseases and MDD.

According to the "hygiene hypothesis" or "Old Friends hypothesis", exposure to certain infectious agents, such as helminths and saprophytic mycobacteria, is important for optimal regulation of the immune system. In modern societies characterized by high levels of hygiene and sanitation, exposure to these organisms is reduced or absent, leading to a loss of immune regulation and tolerance, leading to increased basal cytokine levels and exaggerated immune responses to antigens. This is associated with an increased risk of allergic diseases (such as asthma), immuneinflammatory disorders (such as inflammatory bowel disease), and MDD [30,31]. A related but distinct hypothesis, proposed by Raison and Miller, is known as PATHOS-D (Pathogen Host Defense) [32,33]. PATHOS-D highlights the fact that genetic risk alleles for depression are associated with immune responses to infection, including those involved in the induction of sickness behavior. In this model, the role of exposure or non-exposure to specific pathogens is not central. Instead, emphasis is placed on the fact that genetic variants that were adaptive in defenses against infectious disease (in "ancestral" environments) can be associated with increased immune-inflammatory activity in response to social stressors, leading to the maladaptive phenotype of MDD in "modern" environments [32,33]. This hypothesis is more biologically plausible than the earlier "Old Friends" model [22,34], and has the advantage of being consistent with existing evidence on the relationships between genotype, stressors, and vulnerability towards MDD [23,24,35-37]. However, the two models are not mutually exclusive. It is possible that evolutionarily conserved genetic vulnerability and reduced exposure to microbial signals both contribute in their own way, either additively or synergistically, to the risk of MDD, particularly in the presence of psychosocial risk factors [38–40].

Apart from these hypotheses, there is evidence linking specific infectious diseases to an elevated prospective risk of MDD. For example, the onset of MDD following influenza ("post-influenzal depression") has been documented anecdotally for several decades [41–43], and a prospective association between influenza and subsequent MDD was recently confirmed in a large population-based study [44]. Similar findings have been reported in patients following recovery from other viral

respiratory infections, including SARS [45] and COVID-19 [46,47]. Chronic gum infection (periodontitis) has also been associated with an increased risk of depression [48]. Certain tropical parasitic diseases with an affinity for the central nervous system (CNS), such as Chagas disease and cysticercosis, have also been associated with subsequent MDD. In these cases, such symptoms are likely due to the sequelae of direct brain infection or inflammation induced by the parasite [49]. However, peripheral or systemic mechanisms cannot be entirely ruled out [50], and parasitic diseases such as malaria have been associated with subsequent MDD even in the absence of CNS involvement [51,52]. More recently, attention has also been focused on the relationship between gut microbiota, which can influence immune-inflammatory and stress responses via the gut-brain axis, and MDD. This work provides another potential link between changes in microbial exposure and the risk of MDD [53].

Napolioni and MacMurray have suggested that in populations with high levels of endemic infectious diseases, pathogen pressure may have favored the selection of a specific "high-activity" (G) allele of functional polymorphism (rs1800795) of the interleukin-6 gene (IL6). While this variant is adaptive in defense against microorganisms, it can also be associated with increased CNS inflammation and cognitive impairment [54]. Though these authors did not specifically explore the implications of their model regarding depression, it is notable that several researchers have independently identified links between rs1800795 single-nucleotide polymorphisms (SNPs) and the risk of MDD in diverse populations [23,24,55,56]. In three of these studies, the vulnerability for allele was the "low-activity" (C) allele, whereas in the fourth, homozygotes for both C/C and G/G were at a higher risk of depression than C/G heterozygotes. In these studies, the presence of a specific rs1800795 SNP did not predict depression in isolation, but in interaction with other vulnerability genes or psychological stressors. As in the case of the PATHOS-D and "Old Friends" hypotheses, the picture that emerges is one of complex interactions between genetic vulnerability, exposure to specific micro-organisms, and psychological or social stress, leading to aberrant immuneinflammatory activation that correlates with the symptoms of MDD.

An important limitation of these hypotheses is that they focus largely on MDD and not on BD, though these disorders are closely linked. From an evolutionary perspective, some aspects of BD may have emerged as a natural defense or "override mechanism" to protect individuals and groups from the negative effects of prolonged depression [57]. Building on this perspective, hypomanic "defenses" may have been particularly effective in those ancestral environments where the burden of infectious disease was relatively low. In these environments, prolonged "sickness behaviour"-like phenotypes may have been maladaptive and interfered with other behaviours essential for survival, such as foraging for food, social interactions and reproduction [58]. BD is also associated with elevated levels of peripheral inflammatory markers, and some of these elevations are comparable to those seen in MDD [59]. Thus, any model that attempts to link infectious diseases and MDD should also take BD into consideration.

In the context of these complex and competing hypotheses, it is helpful to examine available data on the relationships between specific infectious diseases and the epidemiology of MDD. The Global Burden of Disease (GBD) studies, conducted over the period 1990-2019, provide estimates of the incidence, prevalence, and disability associated with a wide range of diseases across all the countries and regions of the world [60]. Examination of GBD data has revealed a reduction in the burden of infectious diseases over the past two decades, with a contemporaneous increase in the burden associated with depression [3,61]. This phenomenon has been referred to as an "epidemiological transition," but has not specifically been evaluated in the context of hypotheses such as PATHOS-D or "Old Friends" [62]. Examining the associations between time trends in the occurrence of each of these groups of disorders could help refute, confirm or refine existing theoretical models of the links between infectious diseases and MDD, and yield potential insights into the links between infectious diseases and BD.

The aim of the current study is to examine associations between the incidence of infectious diseases, both as a group and in specific categories, and the incidence of MDD and BD over the past thirty years, based on GBD data. This study is exploratory in nature and is not specifically based on

any of the aforementioned models. However, its results will be examined in terms of how well they align with each model.

# 2. Materials and Methods

The current study was a country-level association study employing both cross-sectional and longitudinal analyses. The estimates of disease incidence provided by the Global Burden of Disease studies, 1990-2019, constituted the primary data source for these analyses. The methodology of the GBD studies has been described in detail by the researchers, and involves collection of data from a wide range of sources, including government and hospital records, registries for specific diseases, and published data in journals, books or online databases. This information is subjected to a rigorous process of cleaning, aggregation and modeling to yield estimates of disease incidence, prevalence and disability-adjusted life years (DALYs) for each country and region [63]. GBD data is freely available for use for academic or research purposes, and can be obtained through database queries from the website of the Institute for Health Metrics and Evaluation (IHME), which is the primary coordinating agency for the GBD studies [64].

# 2.1. Data sources

Information on the incidence of infectious diseases and MDD for the years 1990 and 2019 was obtained from the GBD database [65]. Incidence was selected as the measure of interest to minimize any possible confounding effect arising from chronic or recurrent cases of MDD or BD [66,67]. To reduce the possibility of a confounding effect of life expectancy or age distributions across countries [68], only age-standardized estimates of incidence were retrieved for each category.

For infectious diseases, age-standardized estimates were obtained for the following categories: "upper respiratory infections", "lower respiratory infections", "enteric infections", "neglected tropical diseases and malaria", "intestinal nematode infections", and "other infectious diseases". These categories, which were provided by the GBD database, were selected based on existing theoretical and empirical evidence, as detailed in the Introduction. Data on MDD, BD, and infectious diseases was available for 204 countries.

# 2.2. Confounding or interacting factors

Cross-national variations in the epidemiology of MDD and BD may be due to a variety of factors, which may operate either independent of infectious-immune-inflammatory mechanisms or synergize with them. These factors include climate [69], urbanization [70], and economic factors such as overall economic development and income inequality [71]. Some of these factors may also influence the epidemiology of infectious diseases: for example, tropical and related diseases are more common in countries located close to the equator [72], while higher levels of human development may be associated with lower levels of intestinal infections [73]. To address the possibility of such interactions or confounding effects, data on the following variables was also collected:

- The Human Development Index (HDI), a composite measure of life expectancy, education, and per capita income, as obtained from official United Nations publications [74,75]. This information was available for 124 countries in 1990 and for 187 countries in 2019.
- Levels of urbanization in each country, as measured by the percentage of the total population residing in urban areas, provided by the World Bank's database [76]. This information was available for 200 countries at both time points.
- The Gini coefficient, a measure of income inequality at a national level, from the World Bank's database [77]. This information was available for 168 countries at both time points.
- The average distance of each country from the equator, a proxy measure of climate that has been used in prior epidemiological studies of mental disorders [78,79]. This information was available for all the 204 countries included in this study.

All study variables were tested for normality using the Shapiro-Wilk test. As none of the study variables conformed to a normal distribution, non-parametric statistical tests were used for all further analyses.

In the cross-sectional analyses, Spearman's correlation coefficient was computed to examine the strength and direction of possible monotonic associations between the incidence of each type of infectious disease and the incidence of MDD and BD. To test for the possible confounding or interacting effects of socioeconomic variables and climates, correlations between each of these variables and the prevalence of infectious diseases, MDD and BD were also calculated. Bonferroni's correction for multiple comparisons was applied for all these analyses. If any socioeconomic or climatic variable was significantly associated with either infectious diseases or mood disorders, Spearman's partial correlations were computed. This was done to assess if any associations between infectious diseases and mood disorders remained significant when correcting for these variables.

For the longitudinal analyses, three methods were used. First, cross-lagged correlation coefficients were computed to test for the possibility of a causal relationship between changes in the incidence of infectious diseases and changes in the incidence of mood disorders [80]. Second, the percentage of change of each disorder for the period 1990-2019 was computed, and correlations between these measures of change were computed as for the cross-sectional analyses. To rule out any possible confounding or interacting effects, these tests were also carried out for socioeconomic variables and climate. Third, the chi-square test was used to compare the directions of change in the incidence of mood disorders based on changes in the incidence of each type of infectious disease.

All statistical tests were two-tailed, and a p value of < .05 was used as the threshold for statistical significance. When assessing the strengths of bivariate and partial correlations, standard guideline values for biomedical research were used as follows:  $\varrho \ge 0.8$ , strong correlation;  $\varrho = 0.6$ -0.79, moderate correlation;  $\varrho = 0.3 - 0.59$ , fair correlation;  $\varrho = 0.1 - 0.29$ , weak correlation [81].

# 3. Results

# 3.1. Incidence of infectious diseases and mood disorders in 204 countries, 1990-2019

Data on the incidence of infectious diseases and mood disorders was available for a total of 204 countries. This data, as well as the available data on confounding factors, is summarized in **Table 1**. The incidence of MDD, reported as the median and inter-quartile range (IQR) per 1000 population, was 7.48 (3.05) in 1990 and 7.10 (3.09) in 2019. The incidence of BD was 0.10 (0.04) at both time points.

**Table 1.** Descriptive statistics and changes over time for the incidence of mood disorders and infectious diseases.

Incidence	1990	2019	Change (%)	Significance
Major depressive disorder	7.48 (3.04)	7.10 (3.09)	-3.72 (10.56)	W = 15547.5, $p < .001$
Bipolar disorder	0.10 (0.04)	0.10 (0.04)	0 (6.26)	W = 9039.5, p = .464
Upper respiratory infections	42.01 (9.28)	41.52 (8.51)	-0.80 (5.25)	W = 13013.5, p = .002
Lower respiratory infections	1.59 (0.76)	1.14 (0.54)	-26.75 (12.13)	W = 20903.0, p < .001
Enteric infections	16.19 (7.49)	19.08 (8.32)	12.49 (24.99)	W = 3686.5, p < .001
Intestinal nematode infections	12.93 (33.16)	7.28 (14.22)	-33.76 (54.66)	W = 10982.0, p < .001
Tropical infectious diseases	184.03 (1127.66)	161.69 (812.85)	-20.70 (49.06)	W = 13051.5, p < .001

Other infectious diseases	1 22 (0.90)	1 01 (0 20)	21 77 (12 50)	W = 20904.0,
Other infectious diseases	1.33 (0.89)	1.01 (0.28)	-21.77 (13.58)	<i>p</i> < .001

**Note:** All incidence rates for 1990 and 2019 are given as median (inter-quartile range). **Abbreviations:** W, Wilcoxon's signed-rank sum test statistic; *p*, significance level (*p*-value).

When examining changes in the incidence of each disorder with time, it was noted that the incidence of MDD decreased significantly over the period 1990-2019 (Wilcoxon's W = 15547.5; p < .001), with a rank-biserial correlation of 0.50, indicating a medium effect size. The incidence of BD did not change significantly in either direction during this period (W = 9039.5, p = .464). For infectious diseases, there were significant decreases in the incidence of all categories of disease, except for enteric infections, which showed a significantly large increase over this period (W = 3686.5; p < .001; rank-biserial correlation = -0.65).

## 3.2. Cross-sectional associations between the incidence of infectious diseases and mood disorders

As the first step of the data analysis, cross-sectional correlations between the incidence of each category of infectious diseases and the incidence of both types of mood disorder were calculated for the years 1990 and 2019. These correlations are presented in **Table 2**. It can be observed that in the year 1990, there were no significant correlations between the incidence of MDD and any of the categories of infectious disease. However, the incidence of BD was positively correlated with the incidence of upper respiratory infections, and negatively correlated with every other category of infectious disease. These correlations remained significant after applying Bonferroni's correction for an 8 x 8 table, but were only weak to fair in strength (Q = -0.29 to -0.47). A very similar pattern was obtained when analyzing incidence data for the year 2019; MDD was not correlated with any infectious disease category, while BD remained positively correlated with upper respiratory infections and negatively correlated with other infectious diseases. Scatter plots of all these correlations are provided in **Supplementary Figure 1**.

**Table 2.** Cross-sectional correlations between the incidence of mood disorders and infectious diseases, 1990 and 2019.

Wood Mood				Infectio	ous diseases		
Year	disorder	URI	LRI	Enteric	Nematode	Tropical	Other
1000	MDD	02 (.814)	.07 (.323)	05 (.456)	02 (.814)	03 (.682)	.03 (.711)
1990	BD	.40 (<.001)**	31 (<.001)**	47 (<.001)**	35 (<.001)**	29 (<.001)**	44 (<.001)**
2010	MDD	05 (.525)	.11 (.115)	.07 (.309)	.06 (.362)	.02 (.747)	.09 (.214)
2019 -	BD	.41 (<.001)**	35 (<.001)**	46 (<.001)**	33 (<.001)**	37 (<.001)**	43 (<.001)**

**Note**: All correlations are provided as Spearman's  $\varrho$  (p – value). \* Significant at p < .05, uncorrected. \*\* Significant at p < .05 after Bonferroni's correction for multiple comparisons. **Abbreviations**: MDD, major depressive disorder; BD, bipolar disorder; URI, upper respiratory infections; LRI, lower respiratory infections.

Correlations involving possible confounding or interacting variables are presented in **Supplementary Table 1**. For the year 1990, there was a weak positive correlation between the incidence of MDD and distance from the equator, while the incidence of BD was positively correlated with urbanization and the HDI. The same pattern of associations was observed for 2019, except that the incidence of BD in 2019 was now positively correlated with distance from the equator. Only the associations between BD, urbanization and the HDI remained significant after using the Bonferroni correction. Among infectious diseases, the incidence of upper respiratory infections was positively correlated with HDI and urbanization, and negatively correlated with distance from the equator and the Gini coefficient. The reverse pattern (negative correlations with HDI and urbanization, and positive correlations with the Gini coefficient and distance from the equator) was observed for all other groups of infectious diseases.

As these findings suggested possible confounding effects, partial correlations were computed for all the associations presented in **Table 2**, correcting for all four of these variables. These results

are presented in **Table 3**. It can be noted that after adjusting for economic variables and climate, the incidence of MDD was weakly and negatively correlated with the incidence of enteric infections, tropical diseases, and other infectious diseases in 1990, but not in 2019. The association between BD and upper respiratory infections was no longer significant after these adjustments, but the incidence of BD remained negatively correlated with intestinal nematode infections, tropical diseases, and other infectious diseases both in 1990 and in 2019.

**Table 3.** Partial correlations between the incidence of mood disorders and infectious diseases, 1990 and 2019, adjusted for possible confounding factors.

Year	Mood		Infectious diseases					
Tear	disorder	URI	LRI	Enteric	Nematode	Tropical	Other	
1990 —	MDD	.12 (.209)	04 (.670)	20 (.036)*	16 (.094)	25 (.009)*	26 (.006)*	
	BD	.00 (.963)	.03 (.735)	07 (.481)	31 (<.001)**	21 (.026)*	42 (<.001)**	
2019 —	MDD	.04 (.650)	09 (.269)	04 (.607)	20 (.013)*	14 (.074)	08 (.307)	
	BD	.11 (.158)	18 (.026)*	23 (.003)*	17 (.033)*	29 (<.001)**	30 (<.001)**	

Note: All correlations are provided as Spearman's partial  $\varrho$  (p – value). All correlations are adjusted for Human Development Index (HDI), Gini coefficient, percentage of population residing in urban areas, and distance from the equator. Full descriptions of these variables are given in Section 2.2. \* Significant at p < .05, uncorrected. \*\* Significant at p < .05 after Bonferroni's correction for multiple comparisons. **Abbreviations:** MDD, major depressive disorder; BD, bipolar disorder; URI, upper respiratory infections; LRI, lower respiratory infections.

# 3.3. Longitudinal analyses

# 3.3.1. Cross-lagged regression analyses

To test for the possibility of a causal relationship between changes in the incidence of nfectious diseases and changes in the incidence of mood disorders, cross-lagged regression analyses were carried out for each category of infectious disease. The initial analysis included all 204 countries. However, the accuracy of this analysis was negatively affected by the fact that 68 of these countries had values of zero or close to zero for the incidence of some infectious diseases. To address this limitation, a second analysis was carried out with these countries excluded. These results are presented in **Tables 4a** and **4b**. In these analyses, no evidence was found for a significant causal relationship – in either direction – between any category of infectious disease and mood disorders. A weak trend (p < .2) was found for a possible causal association between LRI and MDD in both analyses.

**Table 4a.** Cross-lagged regression analyses of the relationship between mood disorders and infectious diseases, 1990-2019.

Infectious disease category	Cross-correlation (infectious disease y 1990 x mood disorder 2019)	Cross-correlation (mood disorder 1990 x infectious disease 2019)	Cross-lagged regression coefficient	Significance level
URI				_
x MDD	08	05	029	.681
x BD	34	32	.018	.798
LRI				
x MDD	.12	.03	.097	.168
x BD	34	34	.002	.977
Enteric				_
x MDD	.01	.01	.001	.989
x BD	43	44	.008	.910

Nematode				
x MDD	.09	02	.109	.121
x BD	38	36	022	.755
Tropical				
x MDD	.22	.14	.075	.286
x BD	23	27	.038	.589
Other				
x MDD	.11	.04	.065	.356
x BD	43	39	037	.599

**Table 4b.** Cross-lagged regression analyses of the relationship between mood disorders and infectious diseases, 1990-2019, with outliers excluded.

Infectious disease category	1990 x mood disorder		Cross-lagged regression coefficient	Significance level
	2019)	2019)		
URI				
x MDD	26	22	040	.644
x BD	.14	.14	001	.991
LRI				
x MDD	.20	.07	.133	.123
x BD	20	26	.056	.517
Enteric				
x MDD	.16	.14	.022	.799
x BD	31	29	017	.844
Nematode				
x MDD	01	02	.006	.945
x BD	34	34	006	.945
Tropical				
x MDD	.27	.22	.048	.579
x BD	19	24	.056	.517
Other				
x MDD	.20	.16	.033	.703
x BD	38	34	041	.636

**Note:** The analysis in Table 4b includes 136 countries. 68 countries were excluded due to zero or near-zero incidence estimates for certain infectious disease categories. **Abbreviations:** MDD, major depressive disorder; BD, bipolar disorder; URI, upper respiratory infections; LRI, lower respiratory infections.

# 3.3.2. Relationships between changes in the incidence of infectious diseases and mood disorders over time

Over the period 1990-2019, the incidence of MDD decreased by a median of -3.72% (IQR 10.56; range -35.16% to 34.32%), while the median percentage change for the incidence of BD was zero (IQR 6.26; range -11.54% to 17.24%). Correlations between the percentage changes in the incidence of MDD and BD, and the corresponding changes in the incidence of infectious diseases, for the period 1990-2019 are presented in Table 5. Changes in the incidence of MDD were positively correlated with changes in both upper and lower respiratory infections, and negatively correlated with changes in enteric, nematode, and tropical infections. However, all these correlations were weak ( $\varrho < 0.3$ ) and did not remain significant after correction for multiple comparisons, except in the case of lower respiratory infection. In contrast, changes in the incidence of BD were strongly positively correlated with changes in upper respiratory infections ( $\varrho = .92$ , p < .001) and strongly negatively correlated with

changes in enteric infections ( $\varrho$  = -.82, p < .001). Both of these associations remained significant when correcting for multiple comparisons.

**Table 5.** Correlations between percentage changes in the incidence of infectious diseases and mood disorders, 1990-2019.

Diagnostic category	URI	LRI	Enteric	Nematode	Tropical	Other
MDD	.17 (.013)*	.29 (<.001)**	17 (.013)*	16 (.019)*	21 (.002)*	.13 (.066)
MDD (adjusted)	.06 (.423)	.27 (<.001)**	11 (.176)	11 (.154)	16 (.045)*	.13 (.108)
BD	.92 (<.001)**	.05 (.451)	82 (<.001)**	01 (.893)	29 (<.001)**	12 (.084)
BD (adjusted)	.88 (<.001)**	.00 (.962)	78 (<.001)**	.12 (.125)	23 (.003)*	08 (.289)

**Abbreviations:** MDD, major depressive disorder; BD, bipolar disorder; URI, upper respiratory infections; LRI, lower respiratory infections. **Note:** All correlations are provided as Spearman's  $\varrho$  (p – value). Adjusted correlations are partial correlations adjusted for percentage changes in Gini coefficient and percentage of population residing in urban areas. \* Significant at p < .05, uncorrected. \*\* Significant at p < .05 after Bonferroni's correction for multiple comparisons.

When these correlations were adjusted for changes in urbanization and economic inequality over time, the basic pattern of associations remained unchanged, and changes in the incidence of BD remained strongly correlated with changes in upper respiratory (Q = .88, p < .01) and enteric (Q = .78, p < .01) infections.

# 3.3.3. Categorical associations between changes in the incidence of infectious diseases and mood disorders over time

Comparisons of changes in the incidence of each mood disorder against changes in infectious diseases were also performed using these variables. Complete details of these analyses are presented in Table 6. In these analyses, increases in the incidence of MDD were more common in countries with a corresponding increase in upper respiratory infections over the study period (36/84 vs. 29/120;  $\chi^2$  = 7.95, p = .005). No other significant association was observed for MDD. Increases in the incidence of BD were more common in countries with an increase in upper respiratory infections (78/84 vs 18/120;  $\chi^2$  = 120.22, p < .001), a decrease in enteric infections (50/55 vs 46/149;  $\chi^2$  = 58.12, p < .001), or a decrease in tropical infections (82/157 vs 14/47;  $\chi^2$  = 7.31, p = .007).

**Table 6a.** Categorical associations between changes in the incidence of infectious diseases and changes in the incidence of major depression.

Change in the	Change in the incidence of major				
incidence of	depre	depression		Significance level	
infectious disease	Decreased	Increased			
URI					
Decreased	91 (65.5%)	29 (44.6%)	7.95	.005*	
Increased	48 (34.5%)	36 (55.4%)	7.93	.005	
LRI					
Decreased	138 (99.3%)	64 (98.5%)	0.31	F27+	
Increased	1 (0.7%)	1 (1.5%)	0.31	.537+	
Enteric					
Decreased	33 (23.7%)	22 (33.8%)	2.20	120	
Increased	106 (76.3%)	43 (66.2%)	2.30	.130	
Nematode			0.01	.999+	

Decreased	135 (97.1%)	63 (96.9%)			
Increased	4 (2.9%)	2 (3.1%)			
Tropical					
Decreased	103 (74.1%)	54 (83.1%)	2.01	15/	
Increased	36 (25.9%)	11 (16.9%)	2.01	.156	
Other					
Decreased	138 (99.3%)	63 (96.9%)	1.70	220±	
Increased	1 (0.7%)	2 (3.1%)	1.70	.239†	

**Table 6b.** Categorical associations between changes in the incidence of infectious diseases and changes in the incidence of bipolar disorder.

Change in the					
incidence of			$\chi^2$	Significance level	
infectious disease	Decreased	Increased			
URI					
Decreased	102 (94.4%)	18 (18.8%)	120.22	< 001*	
Increased	6 (5.6%)	78 (81.3%)	120.22	<.001*	
LRI					
Decreased	107 (99.1%)	95 (99.0%)	0.01	000+	
Increased	1 (0.9%)	1 (1.0%)	0.01	.999†	
Enteric					
Decreased	5 (4.6%)	50 (52.1%)	58.12	<.001*	
Increased	103 (95.4%)	46 (47.9%)	36.12	<.001	
Nematode					
Decreased	105 (97.2%)	93 (96.9%)	0.02	.999 <sup>+</sup>	
Increased	3 (2.8%)	3 (3.1%)	0.02	.999'	
Tropical					
Decreased	75 (69.4%)	82 (85.4%)	7.31	007*	
Increased	33 (30.6%)	14 (14.6%)	7.31	.007*	
Other					
Decreased	106 (98.1%)	95 (99.0%)	0.23	.999†	
Increased	2 (1.9%)	1 (1.0%)	0.23	.799'	

Abbreviations: MDD, major depressive disorder; BD, bipolar disorder; URI, upper respiratory infections; LRI, lower respiratory infections. \* Significant at p < .05. † Fisher's exact test.

# 3.4. Subgroup analyses

# 3.4.1. Analyses of tropical and nematode infections

An examination of the incidence data used in this study shows that several countries had estimated incidence rates of zero or close to zero for two categories of infectious diseases – intestinal nematode infections and tropical infections. As the presence of these values could have affected the validity and precision of the pertinent bivariate analyses, sub-group analyses were carried out for these two groups of infections after removing all cases with a negligible estimated incidence.

In the case of intestinal nematode infections, data on 146 countries was analyzed. It was found that the incidence of these infections was not significantly correlated with the incidence of MDD, but remained negatively correlated with the incidence of BD (1990:  $\varrho$  = -.44, p < .001; 2019:  $\varrho$  = -.36, p < .001); this association remained significant even after correction for multiple comparisons and adjustment for possible confounders. Cross-lagged panel analyses did not reveal evidence of a causal association between nematode infections either mood disorder. Correlations of percentage changes showed that changes in nematode infections from 1990 to 2019 were positively correlated with changes in MDD ( $\varrho$  = .36, p < .001) and negatively correlated with changes in BD ( $\varrho$  = -.39, p < .001).

These correlations were attenuated after correcting for changes in economic variables, but remained significant (MDD: Q = .28, p = .002; BD: Q = .24, p = .008).

In the case of tropical infectious diseases, data on 144 countries was analyzed. The incidence of tropical infections was not significantly correlated with the incidence of MDD, but was negatively correlated with the incidence of BD (1990:  $\varrho$  = -.36, p < .001; 2019:  $\varrho$  = -.45, p < .001). After adjusting for confounders, the association with BD remained significant only in 2019 ( $\varrho$  = -.35, p < .001). Crosslagged panel analyses did not reveal possible causal associations for any disorder. Correlations of percentage changes showed that changes in tropical infections from 1990 to 2019 were negatively correlated with changes in the incidence of both MDD ( $\varrho$  = -.23, p = .005) and BD ( $\varrho$  = -.43, p < .001). However, after adjusting for economic variables, only the association with BD remained significant ( $\varrho$  = -.36, p < .001).

# 3.4.2. Analyses of possible interactions between groups of infectious diseases

When analyzing the data on the incidence of specific subgroups of infectious diseases, it was found that there were several significant correlations between them, as can be noted in Supplementary Table 2. It can be observed that the incidence of upper respiratory infections is moderately to strongly negatively correlated with the incidence of all the other infectious disease subgroups (Q = -.44 to -.91, p < .001), while the incidence of other groups of infectious diseases show positive correlations with each other.

In the light of this finding, the analyses reported in sections 3.2 and 3.3.2 were repeated while controlling for the incidence (or change in incidence) of other groups of infectious diseases. In the adjusted cross-sectional analyses, it was found that in the year 1990, the incidence of MDD was negatively correlated with enteric infections ( $\rho = -.16$ , p = .026). The incidence of BD was positively correlated with upper respiratory infections (q = .45, p < .001) and tropical infections (q = .18, p = .013), and negatively correlated with enteric infections (q = -.25, p < .001). However, in the year 2019, only a single significant adjusted correlation was observed: the incidence of BD was negatively correlated with enteric infections (q = -.15, p = .039). All these correlations were weak to fair in strength. In the adjusted longitudinal analyses, changes in the incidence of BD were positively correlated with changes in upper respiratory infections ( $\varrho$  = .66, p < .001) and negatively correlated with changes in enteric infections ( $\rho = -.23$ , p = .001) and tropical infections ( $\rho = -.14$ , p = .046). Changes in the incidence of MDD were positively correlated with changes in lower respiratory infections ( $\rho = .29$ , p < .001) and negatively correlated with changes in enteric (Q = -.14, p = .047) and tropical (Q = 0.21, p = .003) infections. After corrections for multiple comparisons, only two associations remained significant: a positive correlation between changes in upper respiratory infections and BD, and a positive correlation between changes in lower respiratory infections and MDD. A complete set of these analyses can be found in Supplementary Table 3.

# 4. Discussion

The current study examined cross-sectional and longitudinal associations between the incidence of specific groups of infectious diseases and the incidence of mood disorders, including both major depressive disorder (MDD) and bipolar disorder (BD). The categories of infectious diseases analyzed included upper respiratory infections (URI), lower respiratory infections (LRI), enteric infections, intestinal nematode infections, tropical infectious diseases and other infectious diseases. The "other" category consists of vaccine-preventable diseases such as diphtheria, pertussis, tetanus and rubella (measles) [65]. In the time period covered by this study (1990-2019), spanning thirty years, there was a slight but significant decrease in the incidence of MDD but essentially no change in the incidence of BD. All categories of infectious disease showed a significant decrease in incidence, except for enteric infections where a slight increase was observed (Table 1).

In the cross-sectional analyses, there were no significant correlations between the incidence of MDD and any infectious disease category. However, BD showed a specific pattern characterized by positive correlations with the incidence of URI and negative correlations with the other five infectious disease categories (Table 2). This pattern was stable over time and robust to correction for multiple

comparisons. When correcting for possible confounding factors, no consistent pattern of associations was observed for MDD, but BD remained negatively correlated with the incidence of nematode, tropical and other infectious diseases over time (Table 3). When correcting for inter-correlations between categories of infectious disease, a negative correlation between BD and enteric infections was observed at both time points; associations for MDD were weak or inconsistent.

When examining longitudinal associations, three distinct methods were used: cross-lagged correlation analyses, correlations of percentages changes, and categorical associations. The first of these methods did not yield any evidence of a causal association between infectious diseases and either MDD or BD. The second found evidence of (a) a positive association between changes in LRI and changes in MDD, (b) a positive association between changes in URI and in BD, and (c) a negative association between changes in enteric infections and BD, even after correcting for changes in socioeconomic factors, changes in other infectious disease categories, and multiple comparisons. The third found (a) a positive association between changes in the incidence of URI and changes in the incidence of both MDD and BD and (b) negative associations between changes in the incidence of BD and changes in both enteric and tropical infections.

These results will be discussed under four main headings: comparisons between the current study's results and those of other published research; the differential associations of infectious diseases with each type of mood disorder; the relationship of these results to existing hypotheses such as the hygiene hypothesis or PATHOS-D; and possible mechanisms underlying these associations. Following this, the strengths and limitations of the current study, and the broader implications of its results, will be briefly outlined.

# 4.1. Comparisons with the existing literature

The current study suggests that statistically significant relationships exist between the incidence of specific groups of infectious diseases and the incidence of mood disorders. Though this study is the first to make use of global epidemiological data to investigate this relationship, several other researchers have examined the links between infection and mood disorders using other methods. A study of children and adolescents in Taiwan found that bacterial infections were associated with a 2.5-fold increase in the risk of BD and a 2-fold increase in the risk of MDD, with the strongest associations being reported for streptococcal infections [82]. However, this study's findings were not specific to mood disorders: elevated rates of subsequent diagnoses of autism spectrum disorder, attention-deficit/hyperactivity disorder and obsessive-compulsive disorder were also associated with bacterial infection in this sample. A larger study from the same country, including both children and adults and focusing exclusively on intestinal infections, reported similar results: intestinal infections were associated non-specifically with a wide range of mental disorders, including a 1.5 to 2-fold increase in mood disorder risk [83]. This non-specific pattern of association was also observed in a study from the United States, where childhood infections were associated with an increased risk of MDD, but also of anxiety disorders and oppositional-defiant disorder (ODD) [84].

These relationships are not unidirectional. For example, a third Taiwanese study found a prospective association between a diagnosis of BD and the subsequent risk of both URI and LRI, but this association was not specific to infectious diseases and also involved cardiovascular, cerebrovascular, and non-infective gastrointestinal disorders [85]. A study of hospitalizations among patients with BD and related disorders in the United Kingdom found that this disorder was associated with an increased risk of hospitalization for infectious diseases in general, as well as for non-infectious respiratory, digestive and urinary disorders [86]. A follow-up study of cohort of German patients found that MDD was associated with increased rates of LRI and intestinal infections, with a "dose-response" relationship between the severity of MDD symptoms and the risk of these infections [87].

More recent studies in larger cohorts or biobank samples, using genetic markers or risk scores, have underlined the complexity of the association between infectious diseases and mood disorders. In a Mendelian randomization study of over 300,000 participants in the UK Biobank, the genetic risk score for MDD was significantly associated with enteric infections caused by *Escherichia coli* [88].

Complementary to this result, two studies in a Danish cohort of over 65,000 participants, involving both epidemiological and genetic data, found a positive association between infection load and mood disorders, and also suggested that this association might be genetically mediated in the case of MDD, but not of BD [89,90].

In addition to this research on broad categories of infectious disease, some researchers have examined the links between infections with specific pathogens and mood disorders. For example, a prospective relationship has been documented between Epstein-Barr virus (EBV) infection, causing infectious mononucleosis, and MDD both in the short and long terms [91]. Similar associations with MDD have been reported for influenza virus infections [44] and pulmonary tuberculosis [92].

The results of the current study are partly consistent with these results. In both cases, positive associations between certain types of infectious diseases and mood disorders, particularly respiratory infections, have been documented. However, the exact direction of causality cannot be inferred from our data. The one divergent finding was that of a positive association between enteric infections and both MDD and BD in a Taiwanese sample; in the current study, enteric infections were negatively associated with BD. This may reflect differences in levels of analysis; an association that is negative at the cross-national level may be positive in a specific country or region, due to the confounding effects of both population genetics and environmental factors such as diet or social support [10,22].

# 4.2. Differential associations of infectious diseases with major depression and bipolar disorder

This study's findings suggest that there is a differential association between infectious disease groups and distinct categories of mood disorder. In cross-sectional analyses, significant and consistent correlations were observed only for BD. In longitudinal analyses, disorder-specific patterns of association emerged: MDD was positively associated with LRI, while BD was positively associated with URI and negatively associated with enteric and tropical infections.

These findings differ from those of the cohort studies discussed above, which suggest that MDD is associated with enteric infections and is more strongly linked to infectious diseases than BD. Differences in the levels or units of analysis, as mentioned in the previous section, may account for some of these differences, but it is also possible that specific pathogens may interact with host immune-inflammatory factors and other environmental factors, resulting in different outcomes with reference to mood disorder.

The positive association between MDD and LRI is consistent with earlier studies [44,87,92] and is most likely a two-way relationship. Immune-inflammatory changes associated with specific pathogens can trigger MDD in vulnerable individuals [93–95], while MDD may be associated both with reduced immunity to specific respiratory pathogens [96,97]. It is also possible that MDD may be linked to LRI through shared risk or vulnerability factors, such as substance abuse, socioeconomic disadvantage, comorbid medical conditions such as diabetes mellitus, or exposure to environmental pollutants [98–100]. Sometimes the association is more complex and can even span generations: for example, maternal depression has been associated with an increased risk of LRI in children [101].

Regarding the association between URI and BD, there is prior serological and clinical evidence that certain types of viruses causing URI, such as the influenza viruses and coronaviruses, are associated with mood disorder, but this association is not specific to MDD or BD [44,102]. There are case reports of manic episodes being triggered by infection with these viruses [103–106]. A recent systematic review also concluded that viral infections could serve as a trigger for acute mania, but not for depression; however, the authors of this review relied on case reports, and could not establish a specific relationship with URI [107]. A more recent study failed to find an association between bipolar disorder and the presence of antibodies to seasonal coronaviruses [108].

In contrast to the above findings, there is no specific evidence linking BD to an increased risk of or vulnerability to URI as a whole. Research conducted during the COVID-19 pandemic found that genetic vulnerability to BD was associated with susceptibility to this particular infection [109], and that patients with BD had worse outcomes when infected with SARS-CoV-2 [110]. Therefore, it is possible that such a relationship may exist for certain groups or types of viruses associated with URI, but this remains to be demonstrated.

# 4.3. Relationship between the current results and existing hypotheses

The current study was not designed to specifically test any of the existing hypotheses linking infectious diseases and mood disorders. However, some of the results obtained can be interpreted as either supporting or contradicting existing hypotheses, and as pointing towards improved or modified versions of these hypotheses.

First, the current results do not support a direct "protective" relationship between MDD and infectious diseases. If such a relationship existed, one would expect to find a negative correlation between population levels of MDD and at least some groups of infectious diseases [34]. No such relationship could be consistently demonstrated in the current data set, either in cross-sectional or longitudinal analyses. A modified version of this hypothesis postulates that depression can be triggered by certain infections. Once triggered, specific depressive symptoms such as fatigue and social withdrawal act to facilitate recovery and to minimize the risk of contagion or exposure to further infection [111]. This model cannot be either refuted or supported by the current data, as this would require an analysis of recovery and mortality rates from specific infections rather than incidence or prevalence.

Second, the possibility that at least some episodes of MDD can be triggered by infections, particularly respiratory infections, is supported by the current results in which changes in the incidence of URI and LRI were positively associated with changes in the incidence of MDD. Rantala et al. [112] have suggested that MDD can be subtyped on evolutionary grounds based on specific triggering events such as infection, stress, trauma, bereavement and other physical illnesses. They also highlight the fact that short-term adaptive responses to these events can be maladaptive if prolonged, leading to syndromal MDD. They also suggest that this transition is more likely to happen in the setting of certain lifestyle factors, such as diet and insufficient physical activity, that can cause low-grade inflammatory activity. The PATHOS-D model is similar in many aspects to this model: though it does not specify any subtypes of depression, it is also based on the possibility of an evolutionarily ancient, "adaptive" defence against infection turning "maladaptive" in response to social stressors or environmental circumstances [32]. The PATHOS-D and Rantala et al. models are broadly consistent with the current results. However, neither of them can account for the specific association between respiratory infections and MDD identified in this study, and this relationship requires further exploration.

Third, these results provide mixed support for the "Old Friends" or hygiene hypothesis. If this model was applicable at the cross-national level, one would expect a negative correlation between intestinal nematode infections and MDD, but this was not observed in the current study, though nematode infections were negatively correlated with BD. Findings of the longitudinal analyses also failed to establish a relationship between changes in the incidence of these infections and changes in the incidence of MDD. However, changes in enteric infections were negatively associated with changes in both MDD and BD. Therefore, it is still possible that reduced exposure to enteric pathogens – which could serve as a proxy for exposure to "Old Friends" – is associated with an increased risk of mood disorders, and that this association may be – unexpectedly – stronger in BD than in MDD.

Fourth, the current study highlights an important limitation of all the above models: namely, that they do not consider possible relationships between infectious diseases and bipolar disorder. Though some researchers have highlighted possible associations between both past and recent infections and BD [107,113], these findings have not been integrated into a broader model as has been attempted in the case of MDD. Bowins [57] suggested that milder forms of BD (hypomania) could represent a natural "override mechanism" that guards against the maladaptive aspects of depression. If this notion is placed alongside the PATHOS-D or Rantala et al. models, it is possible that hypomania (and possibly BD) would be more common in environments where the burden of infectious disease was relatively low, and the maladaptive aspects of depression would outweigh its benefits. This hypothesis, though tentative, is supported by the results of the current study, in which the incidence of BD was negatively correlated with all groups of infectious disease with the sole exception of URI. Longitudinal analyses also provide some support for this contention, as decreases

in the incidence of enteric and tropical infections were associated with increases in the incidence of BD. Though the current results should be viewed as provisional, they suggest that it may be useful to expand evolutionary, infection-based models of MDD to incorporate BD, or the entire category of mood disorders in general.

Finally, the above models notwithstanding, it is likely that any relationships between mood disorders and infectious diseases are bidirectional and complex. The cross-lagged panel analyses in our study did not support any direct, linear causal relationship between the two. It is likely that any association between infection and mood disorders may apply only to a subset of cases, may depend crucially on the specific properties of individual pathogens [31], and may be influenced by individual vulnerability factors as well as the social, cultural and economic circumstances [114,115].

## 4.4. Possible causal mechanisms

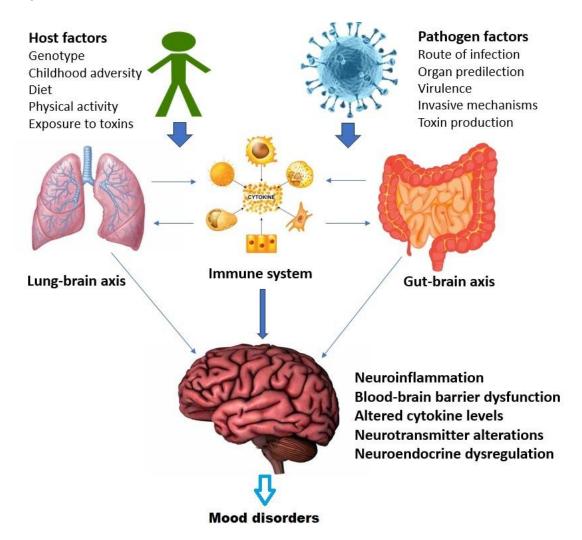
The existence of significant associations between infectious diseases and mood disorders requires an evaluation of the possible mechanistic links between them. The most evident "point of contact" between these groups of disorders is the immune system, or, more broadly speaking, the neuro-immune-endocrine axis. Both MDD and BD are associated with altered levels of circulating cytokines and other inflammatory markers. There are several mechanisms through which these alterations can explain the links between infection and mood disorders. First, some of the immune changes seen in mood disorders may increase susceptibility to infection, as in the case of deficits in T and natural killer (NK) cell activity seen in MDD [97]. Second, certain vulnerability factors for depression, such as childhood adversity, can modify endocrine and immune activity, leading to elevations in "baseline" inflammatory activity. When such vulnerable individuals are exposed to infection, this could trigger an exaggerated or abnormal immune-inflammatory response, leading to an episode of MDD or BD [93,107,115,116]. Third, exposures to infection early in life, during childhood or adolescence, could "prime" the immune system of individuals with genetic variants affecting the expression of inflammatory mediators. This could lead to an exaggerated immune response and symptoms of MDD or BD in response to subsequent stressors, either infective or noninfective in nature [113,117,118]. These explanations are not mutually exclusive, and are each influenced by other risk or protective factors for mood disorders, both genetic and environmental.

More recent research has drawn attention to the role of the microbiota-gut-brain axis in both MDD [119] and BD [120]. In the current study, enteric and nematode infections were negatively associated with the incidence of BD, and decreases in the incidence of enteric infections were associated with increases in both mood disorders, but more specifically with BD. These results are at variance with those of an earlier study, which found that intestinal infections were associated with an increase in subsequent mood disorders [83]. However, it is possible that variations in the incidence of enteric infections may be associated with variations in "non-pathogenic" components of the gut microbiota, which can influence symptomatology and treatment response in BD [121,122]. Alternately, exposure to certain pathogens could induce a state of "immune tolerance", leading to a reduction in the immune-inflammatory dysregulation characteristic of mood disorders [123,124]. This would be a variation of the "hygiene hypothesis" [30,31,39]. A third possibility is that enteric infections could lead patients to receive treatment with antibiotics, which could alter the gut microbiome and trigger depressive or manic episodes [125]. The current results highlight the need to examine the links between enteric infections, gut microbiota and inflammatory profiles in patients with mood disorders.

The current study also found positive associations between rates of respiratory infections and mood disorders, particularly between URI and BD and between LRI and MDD. These results point towards a third avenue that might link infectious and mood disorders, namely the "lung-brain axis". This term refers to bidirectional interactions between the lungs and the brain, mediated through changes in the lung microbiome and the immune system as well as through direct neural links [126,127]. Alterations in the lung microbiome may influence levels of immune and inflammatory activity in the brain, and animal models have shown that infection with specific respiratory pathogens can induce brain inflammation and dysfunction of the blood-brain barrier [128,129].

Similar changes have been implicated in the pathogenesis of MDD and BD [130,131]. It is therefore possible that the results of our study, as well as earlier observations linking viral respiratory infections to mood disorders, could be explained in part by dysregulation of the lung-brain axis.

There is significant cross-talk between these axes at the molecular, cellular and systemic level [132]. This is depicted in Figure 1 above with specific reference to mood disorders. In summary, the relationship between infection and mood disorders may be influenced both by the specific properties of the invading micro-organism and by alterations in the functioning of the lung-brain, gut-brain and immune-endocrine-brain (psychoneuroendocrine) axes. It is unlikely that a simple one-to-one relationship exists between mood disorders and specific infections: rather, this relationship is mediated by a complex interaction between host, pathogen and environmental variables, as well as through alterations in the aforementioned axes [133,134].



**Figure 1.** Interactions between gut-brain, lung-brain and immune systems in the relationship between infection and mood disorders.

# 4.5. Strengths and limitations

The strengths of the current study are its observational, hypothesis-free nature, its use of both cross-sectional and longitudinal methods, its use of the best available data from a large number of countries and territories, and its inclusion of both types of mood disorder. However, there are certain limitations inherent in the methodology adopted. First, as most of the analyses were of an associational nature, it was not possible to draw any definitive conclusions regarding causality. Second, though an attempt was made to assess causality using cross-lagged panel analyses, there are certain limitations inherent in this method [135]. Third, associations identified at the cross-national

or regional level cannot be directly generalized to individuals [136]. Fourth, the relationship between infection and mood disorders is likely to be influenced by several confounding or interacting variables that could not be assessed in the current study, such as variations in allele frequencies of immune-related genes, levels of social support and social capital, or antibiotic prescribing patterns [10,137,138]. Fifth, as infectious diseases were studied as groups, no link could be identified between a specific pathogen (such as the Epstein-Barr virus or *Toxoplasma gondii*) and mood disorders [139,140]. Sixth, some researchers have suggested that links between infectious disease and mood disorders could be influenced by socioeconomic factors [38,39], but this could not be clearly established in the current study despite attempts to control for economic development and inequality. Finally, it is possible that some of the associations between infection and mood disorder could be non-linear in nature, and therefore were not identified using statistical tests based on linear or monotonic models.

## 5. Conclusions

The current study, based on analyses of data from the Global Burden of Disease studies, suggest that significant links exist between specific types of infectious diseases and mood disorders, particularly bipolar disorder. Though subject to the limitations mentioned above, these results provide some support for integrative, evolutionarily-informed models such as PATHOS-D which link mood disorders to evolved defences against infection. The current results also highlight the need for further investigation of the gut-brain and especially the lung-brain axes in the pathogenesis of mood disorders, and to extend models such as PATHOS-D to cover both major depression and bipolar disorder. Though preliminary in nature, this study points towards several promising avenues of future research. These include epidemiological and serological association studies of infectious diseases and mood disorders, cross-national studies examining the links between infection and mood disorders in individual subjects, and studies examining the interactions between infection and other risk or vulnerability factors, such as genetics, childhood adversity, environmental pollutants and stress, in the pathogenesis of mood disorders. Such research could deepen our understanding of the causes of depression and mania, and lead to innovative treatment approaches, such as those targeting gut or lung microbiomes or infection-triggered inflammatory pathways.

**Supplementary Materials:** The following supporting information can be downloaded at the website of this paper posted on Preprints.org, Figure S1: Scatterplots of correlations between the incidence of infectious diseases and mood disorders; Table S1: Correlations involving possible confounding or interacting variables; Table S2: Inter-correlations between the incidence of distinct categories of infectious disease; Table S3: Correlations between the incidence of each category of infectious disease and mood disorders, adjusted for other categories of infectious disease.

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**Data Availability Statement:** The complete dataset used for the purposes of this study will be made available by the author on request.

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