

1 *Review*

2 **Moderate the *MAOA-L* Allele Expression with** 3 **CRISPR/Cas9 System**

4 Martin L. Nelwan
5 Department of Animal Science – Other
6 Nelwan Institution for Human Resource Development
7 Jl. A. Yani No. 24
8 Palu, Sulawesi Tengah, Indonesia
9 Email: mlnelwan2@gmail.com

10 **Abstract:** Antisocial behavior is a behavior disorder inherited according to the inheritance of X-linked
11 chromosome. This disorder derives from mutations in the *MAOA* gene. One of the mutations results in
12 the *MAOA-L* allele activity. The *MAOA-L* allele activity can cause antisocial behavior in both healthy and
13 unhealthy people. Antisocial behavior from healthy males can originate from maltreatment during
14 childhood. Currently, *MAOA* inhibitor can reverse antisocial behavior to normal behavior in animal
15 models. However, this disorder cannot be treated permanently; to treat it permanently in the future,
16 technologies such as CRISPR/Cas9, iPSCs and ssODN are required. These technologies have succeeded to
17 correct erroneous segments in the *F8* gene and *F9* gene. Both genes occupy the X chromosome. The
18 *MAOA* gene also occupies the X chromosome. Therefore, it is reasonable to state that CRISPR/Cas9 and
19 iPSCs technique for instance can be beneficial tools to edit the *MAOA* gene to treat antisocial behavior.
20 CRISPR/Cas9 can be used in combination with iPSCs or ssODN for instance. This combination can
21 greatly help the permanent healing of antisocial behavioral disorders.

22 **Keywords:** advanced therapy; aggressive; antisocial; behavior; *MAOA*

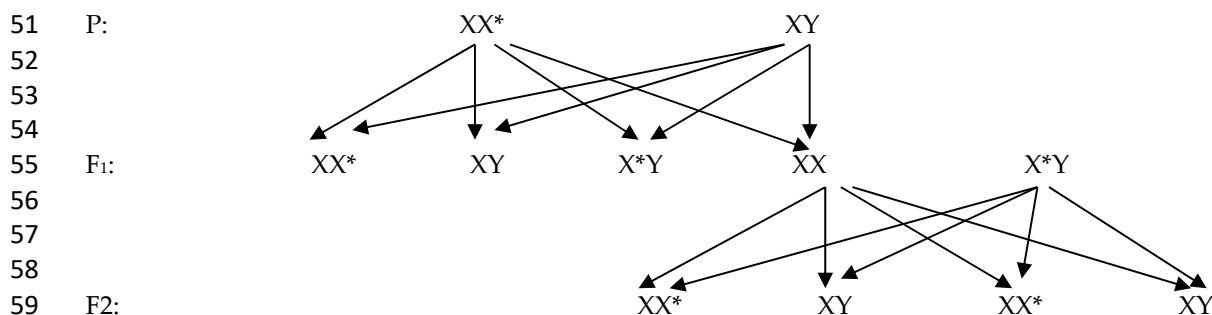
23

24 **1. Introduction**

25 Antisocial behavior is a hereditary disorder inherited through an X-linked recessive inheritance
26 pattern. The *MAOA* gene has correlation with this antisocial behavior [1, 2]. Mutations in this gene result
27 in low *MAOA* (*MAOA-L*) expression [3]. These mutations can create the *MAOA-L* allele. The *MAOA-L*
28 expression affects males nearly entirely and can result in behavior problems such as aggressive and
29 violent eruptions [2, 3]. Antisocial behavior can exist in each family. When parents are carrier female and
30 normal male, $\frac{1}{2}$ female children are carriers and $\frac{1}{2}$ normal female. Male children are $\frac{1}{2}$ normal and $\frac{1}{2}$
31 antisocial behaviors. When parents are normal female and antisocial behavior male, all female children
32 are carrying and all male children are antisocial behaviors (Figure 1). The *MAOA-L* allele is outstandingly
33 general and happens in about 40% [4] or 41% of the Caucasian people [5]. These people have peaceable
34 behavior and have never committed a crime. A study has detected that at least males with this variant
35 had neurobiological framework factors. These factors incline them to violent behavior [3] or antisocial
36 behavior. Maltreatment in children with *MAOA-L* allele can cause antisocial behavior.

37 In animal models, the *MAOA* inhibitor can reverse the antisocial behavior, suggesting that the
38 *MAOA-L* allele expression can be moderated permanently. To reverse antisocial behavior permanently,
39 gene-editing techniques can be used. One of the gene-editing techniques is clustered regularly
40 interspaced short palindromic repeats (CRISPR/Cas9) system. Others are meganucleases (MNs), zinc
41 finger nucleases (ZFNs) and transcription activator-like effectors nucleases (TALENs) [6]. CRISPR/Cas9
42 system gets the benefit of RNA-guided Cas9 nuclease to produce aimed double-stranded DNA breaks
43 (DSBs). Targeted transgenic integration such as homologous recombination (HR) and non-homologous

44 end joining (NHEJ) can correct the DSBs. In addition, the gene-editing technique in combination with
 45 induced pluripotent stem cells (iPSCs) or single-stranded oligonucleotides (ssODN) is useful to fix
 46 erroneous segment in genes such as *MAOA* gene and *F8* gene [6, 7]. For example the CRISPR/Cas9 system
 47 in combination with iPSCs technique has shown its benefit to slow down hemophilia A. Hemophilia A is
 48 inherited according to the X-linked hereditary. Therefore, it is reasonable to state that the iPSCs in
 49 combination with CHRISPR/Cas9 system may be useful to moderate the antisocial behavior in humans,
 50 for instance.



61 Figure 1. In F₁, 50% female is carrier and 50% is normal. 50% male is normal and 50% are affected.
 62 In F₂, all female is carrier and all male is normal.

63 In this article, the author describes progress in the study of antisocial behavior. The author
 64 focuses on the biological aspects and gene therapy. The biological aspects include the *MAOA* gene,
 65 mutations in the *MAOA* gene and antisocial behavior and treatment with gene therapy. Gene therapy
 66 includes CRISPR/Cas9 system in combination with iPSCs or HR-based method.

67 2. Genes in Antisocial Behavior

68 Gene is fundamental bodily and functional unit of heredity. Genes consist of DNAs and supply
 69 instructions to build protein molecules. Changes can occur in a gene and can cause protein destruction. A
 70 gene change is a stable mutation in the DNA. A condition derives from changes in at least one gene stated
 71 as a hereditary disease [8]. For example, hereditary diseases can include such as hemophilia A and
 72 antisocial behavior. Antisocial behavior derived from the mutation of the *MAOA* gene into the *MAOA-L*
 73 allele.

74 2.1. The *MAOA* Gene

75 “Monoamine oxidase A” is the formal name of the *MAOA* gene. *MAOA* is the gene formal
 76 symbol. Other names of these genes include *BRNRS* and *MAO-A*. The gene supplies directive for making
 77 monoamine oxidase A [3]. The *MAOA* gene spans at least 60 kb and consists of 15 exons. The *MAOA* gene
 78 displays the same exon-intron organization. Exon 12 encodes for the covalent FAD-binding site. This
 79 exon is the most conserved exon [9-10]. The *MAOA* gene occupies the p arm of the X chromosome at
 80 position 11.3; Xp11.3 [9]. This gene includes base pairs 43,654,907 to 43,746,824 on the X chromosome [3,
 81 11].

82 The *MAOA* gene is one of two neighboring gene families. The other gene is *MAOB*. The *MAOA*
 83 and *MAOB* derived from duplication of the *MAO* gene [10]. The external mitochondrial membrane
 84 expresses these two genes [9-10]. Chen *et al* stated that these two genes oxidize neurotransmitters and
 85 dietary amines [10]. The regulation of neurotransmitters activity is vital in sustaining standard mental

86 conditions [3, 10]. Chen *et al* localized the *MAOA* and *MAOB* genes within a region of about 240 kb. The
87 *MAOA* gene encodes mitochondrial enzymes and catalyzes the oxidative deamination of amines [9-10].
88 These include such as dopamine, norepinephrine, and serotonin as substrates [3, 10-13]. The *MAOB* gene
89 prefers phenylethylamine as substrates [10, 13].

90 The *MAOA* gene has significant roles in the metabolism of neuroactive and vasoactive amines.
91 This metabolism occurs in the central nervous system and peripheral tissues [3,8]. Ou *et al.*, established
92 that serum hunger-made apoptosis in culture neuronal cell line enhanced demonstration of *MAOA*. In
93 addition, this serum enhanced demonstration of p38 kinase and caspase-3. This apoptosis diminished bd-
94 2 and R1. *MAOA* and R1 were upstream of caspase-3. Both of them were downstream of p38 kinase and
95 *BCL2* in the apoptotic signaling pathway. Moreover, Ou *et al.*, stated that serum starvation of cortical
96 brain cells from *Maoa*-deficient mice resulted in reduced apoptosis contrasted with wild-type (WT) mice.
97 *MAOA* and R1 were involved in the *MYC*-made proliferative signaling pathway in the attendance of
98 serum. Both of them function upstream of cyclin D1 and E2F1 in the cell proliferation pathway. The
99 *MAOA* inhibitor could avoid apoptosis. [10, 14].

100 2.2. Mutations in the *MAOA* Gene for the Antisocial Behavior

101 Antisocial behavior can happen in healthy people [2, 15]. Antisocial behavior is cross-transmitted
102 with other dyscontrol disorders. This behavior has several well-defined biological correlates. These
103 include injured frontal lobe function, leading to reduced ability to control behavior. Aggression is an
104 important manifestation of antisocial behavior. [15]. Caspi *et al* studied male children from birth to
105 adulthood subjected to maltreatment. The authors stated that maltreated children with a genotype
106 conferring the *MAOA-H* allele expression were less likely to develop antisocial behavior. It shows that
107 children with the *MAOA-L* allele expression correspond to antisocial behavior [10, 16]. This means that
108 environment has important role developing antisocial behavior in humans.

109 In 1981, Pinter *et al* allocated the *MAOA* gene on the human X chromosome [5, 10, 17]. Later, the
110 *MAOA-L* allele activity and antisocial behavior in male mice with an X chromosome deletion were linked.
111 In addition, Cases *et al* reported that deletion in the *MAOA* gene in mice revealed an increase of
112 norepinephrine, serotonin and dopamine. Moreover, this mutation raised aggression in male mice [10, 17,
113 18]. Reti *et al* introduced that Caucasians with *MAOA-L* allele had antisocial behavior around 41% [5],
114 supporting a link between *MAOA-L* allele and antisocial behavior.

115 McDermott *et al* conducted a behavioral study in humans to link behavior and environment
116 influence. In this study, the authors paid male subjects to penalize those they considered had stolen
117 money from them. The authors adjusted the amount of money lost from them to their enemies.
118 McDermott *et al* reported that people with *MAOA-L* had violent behavior penalize their enemies. The
119 connection was critical when the quantity of money was higher, suggesting an environmental interaction.
120 It shows that heredity can play a role in the behavior and daily decisions taken [10, 19].

121 Ziemans *et al* stated that a single-nucleotide polymorphism (rs6609257) considerably influenced a
122 brain network activity. This network includes frontal, parietal and occipital areas. The authors indicated
123 that the rs6609257 occupies ~ 6.6 kb downstream of the *MAOA* gene on the human X chromosome.
124 Ziemans *et al* showed that improved activity in this network had correlation with visuospatial working
125 memory (VSWM) capacity in the order predicted externalizing symptoms. The authors indicated that a
126 higher working memory capacity had not associated with fewer externalizing symptoms. However, these
127 externalizing symptoms associated with aggressive/oppositional behavior. In this study, the authors
128 proposed a mediating function or working memory brain activity in connecting the *MAOA* gene to
129 aggressive behavior [20]. Furthermore, Marquez *et al* showed that male rats, which surrendered to
130 pressure-induced experiences through peripubertal, show aggressive behavior at maturity. The authors
131 indicated that treatment with an *MAOA* inhibitor reversed the peripuberty pressure-induced antisocial

132 behaviors. Marquez *et al* showed that biological factors, which are activated through maltreatment, are
133 the cause of antisocial behavior [21]. It suggests that education not to do violent behavior is very
134 important. It can help to reduce “antisocial behavior” among people.

135 Maltreatment in childhood (G x E) can result in emotional and antisocial behavioral problems in
136 youth. These people have low variability on the variable number tandem repeats (VNTR) polymorphism
137 of the MAOA gene [22-23, 25-26]. The VNTR polymorphism in human consists of 30 base pairs in length.
138 These include 2R, 3R, 3.5R, 4R and 5R copies of the repeat series. The polymorphism demonstrated
139 influence on transcriptional activity of the MAOA gene promoter. The 3.5R and 4R repeats are transcribed
140 more effectively than those with 2R and 3R copies [2, 10, 23, 25-26]. Males with a 2R variant have a level
141 of serious criminal behavior and violent behavior. Effects for females are alike, but weaker [10, 25]. The
142 effect 5R is unclear [3]. The 2R promoter displays many inferior levels of promoter activity than the other
143 promoters.

144 Behavior disorder due to abuse has contradictorily established a connection between the MAOA-
145 L allele and antisocial behavior. The MAOA-L allele activity raises the risk of behavior disorder and
146 antisocial behavior traits. This happens to young people who experience maltreatment during childhood
147 [22, 26-29]. In addition, non-linear interactions between the MAOA gene and violence have been found [2,
148 27]. It suggests that people with the MAOA-L allele can be hypocrite once a certain violent level are
149 detected.

150 Three mutations occur in the MAOA-L allele so far. These include nonsense (Brunner syndrome),
151 missense (autism), and a deletion (Norrie disease). Brunner syndrome and autism correspond to
152 aggressive behavior. Brunner syndrome has antisocial behavior and autism has auto-aggressive behavior,
153 while Norrie disease corresponds to autistic-like behavior. Brunner syndrome, autism and Norrie disease
154 belong to intellectual disability (ID). Brunner syndrome shows stress-induced aggressive and violent
155 behavior in addition to borderline ID [30]. This shows that antisocial behavior can include borderline
156 behavior. Furthermore, antisocial behavior can exist both in healthy [2, 4] and in unhealthy people [30].

157 Transgenic mice for antisocial behavior researches can be obtained. Use of mice will be helpful to
158 conduct research for treating disorders inherited through X-linked recessive configurations. These
159 disorders can include such as hemophilia B and antisocial behavior. For example, to diminish antisocial
160 behavior, advanced therapy such as CRISPR/Cas9 system in combination with iPSCs technology can be
161 used. This combination can be helpful to treat diseases inherited through X-linked pattern [6-7].

162 3. The Gene-Editing Technique

163 Currently, four methods are available for targeted integration of transgenic. These include NHEJ,
164 microhomology-mediated end joining (MMEJ), HR, and homology-mediated end joining (HMEJ) (Table 1).
165 The NHEJ-based method presented random directions in integration and various types of indels at the
166 junctions. NHEJ is active in the entire cell cycle [35-36]. The MMEJ-based method displayed low
167 efficiency in cultured cells. MMEJ is active in the early S/G1 phase [35, 37]. The HR-mediated method
168 allows correct insertion of large fragments. This method is commonly inefficient in animal embryos and
169 tissues *in vivo*. HR is active during the late S/G2 phase only. Finally, the HMEJ-based method achieved
170 transgenic integration in mouse and monkey embryos, as well as in hepatocytes and neurons *in vivo* with
171 high efficiency. HMEJ is active in the early S/G1 phase. All of the methods can be useful for generating
172 animal models and for targeted gene therapies [35].

173 3.1. Gene-editing in X-linked Disorders

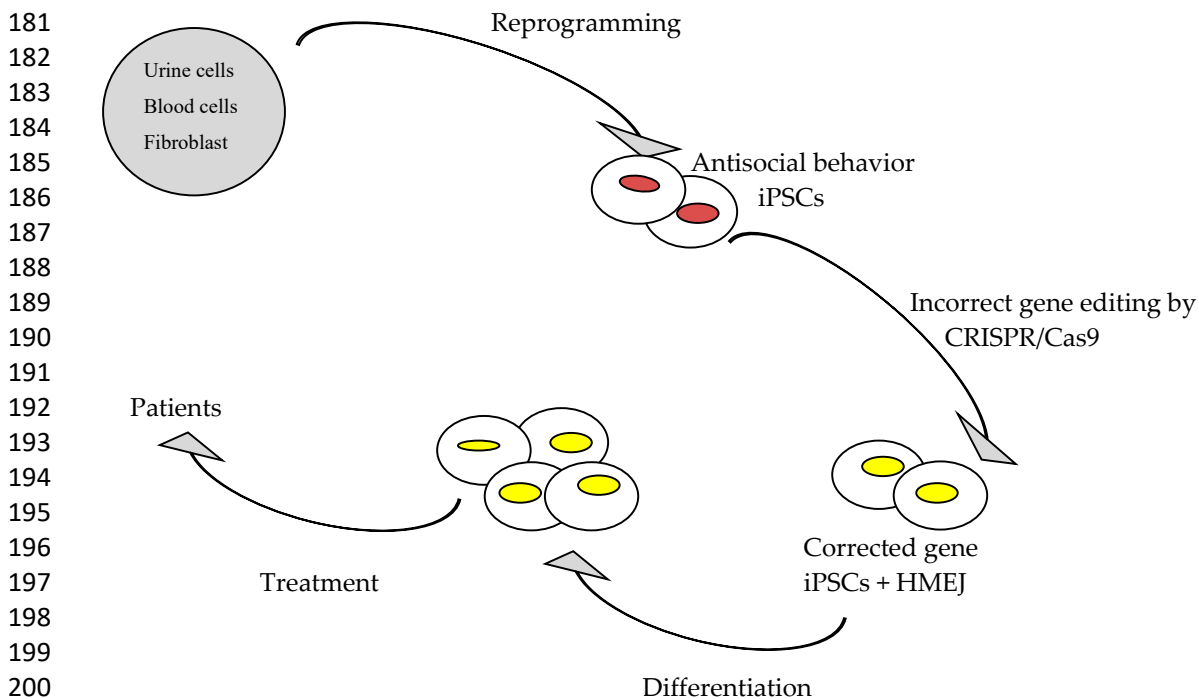
174 The gene-editing method uses creator of nucleases to edit incorrect gene. This method also uses
175 the cellular repair technique to exactly alter incorrect string. For example, to identify the selected genomic

176 location and its transfected into the cell, an artificial chain-precise is designed. It produces DSBs at the
 177 location [31]. The NHEJ [31, 32, 33, 34], MMEJ, HR [34] and HMEJ can repair DSBs. Furthermore,
 178 nucleases such as TALENs and CRISPR/Cas9 systems can induce DSBs in a targeted genomic locus [35].

Table 1. Methods for targeted integration transgenic

Protocol	Advantage/disadvantage	Cell cycle	References
HR	Inefficiency	S/G2	[35]
MMEJ	High efficiency	Early S/G1	[35]
NEHJ	Random directions	entire phase	[35, 36]
HMEJ	Low efficiency	early S/G1	[35, 37]

179
 180



201 **Figure 1.** Treatment of antisocial behavior uses CRISPR/Cas9, iPSCs and HMEJ-based method.

202 The *F8* gene and *F9* gene occupy the X chromosome. It is the same as the *MAOA-L* allele that also
 203 occupies the X chromosome in the chromosome map. Currently, with the development of CRISPR/Cas9
 204 system, mutations in genes *F8* and *F9* can be corrected. For example, the CRISPR/Cas9 in combination
 205 with iPSCs or ssODN can correct erroneous segments in these genes in animal models. CRISPR/Cas9 in
 206 combination with iPSCs is gene-editing technique for hemophilia A. Park *et al* edited mutations in the *F8*
 207 gene with this combination. These authors practically rescued the Factor VIII deficiency in a hemophilia
 208 mouse model [38]. CRISPR/Cas9 in combination with ssODN is gene-editing technique for hemophilia B.
 209 Guan *et al* generate mutated mouse strains for hemophilia B, and then cured these strains *in vivo* by
 210 hydrodynamic tail injection of a plasmid. The plasmid encodes Cas9 and the sgRNA in combination with
 211 ssODN containing the edited string [39]. It seems that these techniques can be potential tools to treat
 212 antisocial behavior.

213 3.2. Gene-editing for Antisocial Behavior

214 Currently, there has not yet been gene therapy to treat antisocial behavior. However, this
215 technique may be a useful tool for treating this disease. To treat antisocial behavior, CRISPR/Cas9 system,
216 iPSCs technique, and targeted transgenic integration such as HR-based method and HMEJ-based method
217 can be used. For example, CRISPR/Cas9 system corrects the incorrect gene in iPSCs and constructs DSBs.
218 Then, HMEJ for instance will correct the nick. It will correct segments or remove the 30-bp copy in the
219 string. This gene-corrected is differentiated into suitable somatic cells. Furthermore, patients can obtain
220 the corrected gene by giving it to patients (Figure 2).

221 To alter the *MAOA-L* allele needs gene-editing technique. In animal models, this technique is a
222 useful tool to treat an X-linked recessive disorder such as hemophilia A and hemophilia B. The
223 CRISPR/Cas9 system in combination with such as the iPSCs technique or HMEJ-based method is useful
224 to fight antisocial behavior.

225 4. Conclusions

226 Antisocial behavior is a violent behavior inherited according to the inheritance of sex-linked
227 recessive allele. A violent behavior derives from mutations in the *MAOA* gene. These mutations can
228 result in, such as autism, Brunner syndrome, and antisocial behavior. The *MAOA-L* allele expression
229 corresponds to antisocial behavior. Environmental factors such as maltreatment can cause antisocial
230 behavior in male children. To treat permanently this behavior, it is impossible at present. In the future, to
231 treat this behavior, a gene-editing tool such as TALENs or CRISPR/Cas9 systems can be used. For
232 example, CRISPR/Cas9 system can correct erroneous segments in the sex-linked disorders. CRISPR/Cas9
233 system can be used in combination with iPSCs technique and HMEJ-based method for instance. For
234 example, CRISPR/Cas9 in combination with iPSCs has corrected erroneous segments in the *F8* gene in the
235 animal models. These findings give hope to treat antisocial behavior with the CRISPR/Cas9 system in
236 combination with iPSCs technique for instance. It can alter the *MAOA-L* allele to normal allele to result in
237 normal behavior. This combination is a promising tool to treat permanently the antisocial behavior in
238 both healthy and unhealthy people.

239 **Acknowledgement:** The author has role in designing of the study and writing of the manuscript.

240 **Author Contributions:** MLN designed the study, performed the literature searches, wrote the first draft of the
241 manuscript and involved in revising the manuscript critically for significant intellectual need. The author read and
242 approved the final manuscript.

243 **Conflict of Interests:** The author declares no conflict of interest.

244 References

- 245 1. Xu, M.K.; Gaysina, D.; Tsonaka, R.; Morin, A.J.S.; Croudace, T.J.; Barnett, J.H.; *et al.* Monoamine Oxidase A
246 (*MAOA*) Gene and Personality Traits from Late Adolescence through Early Adulthood: A Latent Variable
247 Investigation. *Front. Psychol.* **2017**; *8*: 1736, doi:10.3389/psyg.2017.01736.
- 248 2. Kolla, N.J.; Meyer, J.; Sanches, M.; Charbonneau, J. Monoamine Oxidase-A Genetic Variants and Childhood
249 Abuse Predict Impulsiveness in Borderline Personality Disorder. *Clinical Psychopharmacology and Neuroscience*
250 **2017**; *15(4)*: 343-351, doi:10.9758/cpn.2017.15.4.3433.
- 251 3. Genetics Home Reference (2018): *MAOA*, <https://ghr.nlm.nih.gov/gene/MAOA>.
- 252 4. Hunter, P. The Psycho gene. *EMBO reports* **2010**, *11* 9: 667.
- 253 5. Reti, E.M.; Xu, J.Z.; Yanofski, J.; McKibben, J.; Uhart, M.; Cheng Y-J, *et al.* MAO regulates antisocial personality in
254 Caucasians with no history physical abuse. *Compr. Psychiatry* **2011**; *52(2)*: 188-194.

- 255 6. Nelwan, M.L. Hemophilia A and Induced Pluripotent Stem Cells. *Journal of Advances in Biology & Biotechnology*
256 **2017**; 14(3): 1-11, doi:10.9734/JABB/2017/35111.
- 257 7. Nelwan, M.L. Friedreich Ataxia: Treatment with Genetic Approach. *Journal of Advances in Biology & Biotechnology*
258 **2017**; 14(4): 1-12,doi:10.9734/JABB/2017/36113.
- 259 8. Nelwan, M.L. Treat Oculocutaneous Albinism with Gene Therapy. *Journal of Advances in Biology & Biotechnology*
260 **2017**; 16(3): 1-12. DOI:10.9734/ JABB/2017/38504.
- 261 9. Grimsby, J.; Chen, K.; Wang, L.-J.; Lan, N.C.; Shih, J.C. Human Monoamine Oxidase A and B genes exhibit
262 identical exon-intron organization. *Proc. Nat. Acad. Sci.* **1991**; 88: 3637-3641.
- 263 10. Online Mendelian Inheritance in Man (2017): Monoamine Oxidase A; MAOA, <http://omim.org/entry/309850>.
- 264 11. NCBI Gene, <https://www.ncbi.nlm.nih.gov/gene/4128>.
- 265 12. Chen, K.; Holschneider, D.P.; Wu, W.; Rebrin, I.; Shih, J.C. A spontaneous point mutation produces monoamine
266 oxidase A/B knock-out mice with greatly elevated monoamines and anxiety-like behavior. *J. Biol. Chem.* **2004**;
267 279: 39645-39652, doi:10.1074/jbc.M405550200.
- 268 13. UniProt, <http://www.uniprot.org/uniprot/P21397>.
- 269 14. Ou, X.-M.; Chen, K.; Shih, J.C. Monoamine oxidase A and repressor R1 are involved in appropriate signaling
270 pathway. *Proc. Nat. Acad. Sci.* **2006**; 103 (29): 10923-10928, doi:10.1073/pnas.0601515103.
- 271 15. Sjöberg, R.L.; Ducci, F.; Barr, C.S.; Newman, T.K.; Dell'Osso, L.; Virkkunen M, *et al.* Non-Additive Interaction of
272 a Functional MAO-A VNTR and Testosterone Predicts Antisocial Behavior. *Neuropsychopharmacology* **2008**; 33(2):
273 425-430, doi:10.1038/sj.npp.1301417.
- 274 16. Caspi, A.; McClay, J.; Moffitt, T.Y.; Mill, J.; Martin, J.; Craig, I.W.; *et al.* Role of genotype in the cycle of violence in
275 maltreated children. *Science* **2002**; 297: 851-854.
- 276 17. Pintar, J.E.; Barbosa, J.; Francke, U.; Castiglione, C.M.; Hawkins, M.; Breakefield, X.O. Gene for monoamine
277 oxidase type A assigned to the human X chromosome. *The Journal of Neuroscience* **1981**; 1(2): 166-175.
- 278 18. Cases, O.; Seif, I.; Grimsby, J.; Gaspar, P.; Chen, K.; Pournin, S.; *et al.* Aggressive Behavior and Altered Amounts
279 of Brain Serotonin and Norepinephrine in Mice Lacking MAOA. *Science* **1995**; 268(5218): 1763-1766.
- 280 19. McDermott, R.; Tingley, D.; Cowden, J.; Frazzetto, G.; Johnson, D.D.P. Monoamine oxidase A gene (MAOA)
281 predicts behavioral aggression following provocation. *Proc Nat. Acad. Sci.* **2009**; 106(7): 2118-2123,
282 doi:10.1073/pnas.0808378106.
- 283 20. Ziermans, T.; Dumontheil, I.; Roggeman, C.; Peyrard-Janvid, M.; Matsson, H.; Kere, J.; *et al.* Working memory
284 brain activity and capacity link MAOA polymorphism to aggressive behavior during development. *Transl.*
285 *Psychiatry* **2012**; 2 e85.
- 286 21. Marquez, C.; Poirier, G.L.; Cordero, M.I.; Larsen, M.H.; Groner, A.; Marquis, J.; *et al.* Peripuberty stress leads to
287 abnormal aggression, altered amygdale and orbitofrontal reactivity and increased prefrontal MAOA gene
288 expression. *Transl. Psychiatry* **2013**; e216.
- 289 22. Hart, H.; Lim, L.; Mehta, M.A.; Chatzieffraimidou, A.; Curtis, C.; Xu, X.; *et al.* Reduced functional connectivity of
290 fronto-parietal sustained attention networks in severe childhood abuse. *PLoS ONE* **2017**; 12(11): e0188744,
291 10.1371/journal.pone.0188744.
- 292 23. Meyer-Lindenberg, A.; Buckholtz, J.W.; Kolachana, B.; Hariri, A.R.; Pezawas, L.; Blasi, G.; *et al.* Neural
293 mechanism of genetics risk for impulsivity and violence in humans. *PNAS* **2006**; 103(16): 6269-6274,
294 doi:10.1073/pnas.0511311103.
- 295 24. Zhang, Y.; Ming, Q.-s.; Yi J-y, Wang, X.; Chai, Q.-l.; Yao, S.-q. Gene-Gene-Environment Interactions of Serotonin
296 Transporter Monoamine Oxidase and Childhood Maltreatment Predict Aggressive Behavior in Chinese
297 Adolescents. *Front. Behav. Neurosci.* **2017**; 11: 17, doi:10.3389/fnbeh.2017.00017.
- 298 25. Guo, G.; OU, X.-M.; Roettger, M.; Shih, J.C. The VNTR 2 repeat in MAOA and delinquent behavior in adolescence
299 and young adulthood: associations and MAOA promoter activity. *European Journal of Human Genetics* **2008**; 16:
300 626-634, doi:10.1038/sj.ejhg.5201999.
- 301 26. Young, S.E.; Smolen, A.; Hewitt, J.K.; Haberstick, B.C.; Stallings, M.C.; Corley, R.P.; Crowley, T.J. Interaction
302 Between MAO-A Genotype and Maltreatment in the Risk for conduct Disorder: Failure to Confirm in Adolescent
303 Patients. *Am J Psychiatry* **2006**; 163: 1019-1025.
- 304 27. Ouellet-Morin, I.; Côté, S.M.; Vitaro, F.; Hébert, M.; Carbonneau, R.; Lacourse, E.; *et al.* Effects of the MAOA gene
305 and levels of exposure to violence on antisocial outcomes. *The British Journal of Psychiatry* **2016**; 208: 42-48,
306 doi:10.1192/bjp.bp.114.162081.

- 307 28. Godar, S.C.; Fite, P.J.; McFarlin, K.M.; Bortolato, M. The role of monoamine oxidase A in aggression: current
308 translational developments and future challenges. *Prog Neuropsychopharmacol Biol Psychiatry* **2016**; *69*: 90-100.
309 doi:10.1016/j.pnpbp.2016.01.001.
- 310 29. McSwiggan, S.; Elger, B.; Appelbaum, P.S. The Forensic Use of Behavioral Genetics in Criminal Proceedings:
311 Case of the MAOA-L Genotype. *Int J Law Psychiatry* **2017**; *50*: 17-23. doi:10.1016/j.ijlp.2016.09.005.
- 312 30. Piton, A.; Poquet, H.; Redin, C.; Masurel, A.; Laurel, J.; Muller, J.; *et al.* 20 ans après: a second mutation in MAOA
313 identified by targeted high-throughput sequencing in a family with altered behavior and cognition. *European*
314 *Journal of Human Genetics* **2013**; *22*: 776-783. doi:10.1038/ejhg.2013.243.
- 315 31. Ding, Y.; Li, H.; Chen, L.-L.; Xie, K. Recent Advances in Genome Editing Using CRISPR//Cas9. *Frontiers in Plant*
316 *Science* **2016**; Volume 7, Article 703, doi:10.3389/fpls.2016.00703.
- 317 32. Eid, A.; Mahfouz, M.M. Genome editing: the road of CRISPR/Cas9 from bench to clinic. *Experimental and*
318 *Molecular Medicine* **2016**; *46*, e265, doi:10.1038/emm.2016.111.
- 319 33. Chu, V.T.; Weber, T.; Graf, R.; Sommermann, T.; Petsch, K.; Sack, U.; Volchkov, P.; Rajewsky, K.; Kühn, R.
320 Efficient generation of *Rosa26* knock-in mice using CRISPR/Cas9 in C57BL/6 zygotes. *BMC Biotechnology* **2016**;
321 *16*:4, doi:10.1186/s12896-016-0234-4.
- 322 34. Kawahara, A.; Hisano, Y.; Ota, S.; Taimatsu, K Site-Specific Integration Exogenous Genes Using Genome Editing
323 Technologies in Zebrafish. *Int. J. Mol. Sci.* **2016**; *17*: 727. doi:10.3390/ijms17050727.
- 324 35. Yao, X.; Wang, X.; Hu, X.; Liu, Z.; Liu, J.; Zhou, H.; *et al.* Homology-mediated end joining-based targeted
325 integration using CRISPR/Cas9. *Cell Research* **2017**; *27*: 801-814, doi:10.1038/cr.2017.76.
- 326 36. Mao, Z.; Bozzella, M.; Seluanov, A.; Gorbuanova, V. DNA repair by nonhomologous end joining and
327 homologous recombination during cell cycle in human cells. *Cell Cycle* **2009**; *7*(18): 2902-2906.
- 328 37. Nakade, S.; Tsubota, T.; Sakane, Y.; Kume, S.; Sakamoto, M.; Obara, M. Microhomology-mediated end-joining-
329 dependent integration of donor DNA in cells and animals using TALENs and CRISPR/Cas9. *Nature*
330 *Communication* **2014**; *5*: 5560, 10:1038/ncomms6560.
- 331 38. Park, C.Y.; Kim, D.H.; Son, J.S.; Sung, J.J.; Lee, J.; Bae, S.; *et al.* Functional Correction of Large Factor VIII Gene
332 Chromosomal Inversions in Hemophilia A Patient-Derived iPSCs Using CRISPR/Cas9. *Cell Stem Cell* **2015**; *17*(2):
333 213-220.
- 334 39. Guan, Y.; Ma, Y.; Li, Q.; Sun, Z.; Ma, L.; Wu, L.; *et al.* CRISPR/Cas9-mediated somatic correction of a novel
335 coagulation factor IX gene mutation ameliorates hemophilia in mouse. *EMBO Molecular Medicine* **2016**; *8*: 477-488,
336 doi:10.15252/emmm.201506039.