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### **Understanding and predicting suicidality using a combined genomic and clinical risk assessment approach** OPEN

A B Niculescu, D F Levey, P L Phalen, H Le-Niculescu, H D Dainton, N Jain, E Belanger, A James, S George, H Weber, D L Graham, R Schweitzer, T B Ladd, R Learman, E M Niculescu, N P Vanipenta, F N Khan, J Mullen, G Shankar, S Cook, C Humbert, A Ballew, M Yard, T Gelbart, A Shekhar, N J Schork, S M Kurian, G E Sandusky & D R Salomon

doi:10.1038/mp.2015.112

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### **Genetics and intelligence differences: five special findings** OPEN

R Plomin & I J Deary

doi:10.1038/mp.2014.105

[Abstract](#) | [Full Text](#) | [PDF](#)

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### **Phenome-wide analysis of genome-wide polygenic scores** OPEN

E Krapohl, J Euesden, D Zabaneh, J-B Pingault, K Rimfeld, S von Stumm, P S Dale, G Breen, P F O'Reilly & R Plomin

doi:10.1038/mp.2015.126

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### **Subcortical brain alterations in major depressive disorder: findings from the ENIGMA Major Depressive Disorder working group** OPEN

L Schmaal, D J Veltman, T G M van Erp, P G Sämann, T Frodl, N Jahanshad, E Loehrer, H Tiemeier, A Hofman, W J Niessen, M W Vernooij, M A Ikram, K Wittfeld, H J Grabe, A Block, K Hegenscheid, H Völzke, D Hoehn, M Czisch, J Lagopoulos, S N Hatton, I B Hickie, R Goya-Maldonado, B Krämer, O Gruber, B Couvy-Duchesne, M E Rentería, L T Strike, N T Mills, G I de Zubicaray, K L McMahon, S E Medland, N G Martin, N A Gillespie, M J Wright, G B Hall, G M MacQueen, E M Frey, A Carballo, L S van Velzen, M J van Tol, N J van der Wee, I M Veer, H Walter, K Schnell, E Schramm, C Normann, D Schoepf, C Konrad, B Zurowski, T Nickson, A M McIntosh, M Pappmeyer, H C Whalley, J E Sussmann, B R Godlewska, P J Cowen, F H Fischer, M Rose, B W J H Penninx, P M Thompson & D P Hibar

doi:10.1038/mp.2015.69

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### **The 3q29 deletion confers >40-fold increase in risk for schizophrenia** OPEN

J G Mulle

doi:10.1038/mp.2015.76

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### **Whole-genome sequencing suggests a chemokine gene cluster that modifies age at onset in familial Alzheimer's disease** OPEN

M A Lalli, B M Bettcher, M L Arcila, G Garcia, C Guzman, L Madrigal, L Ramirez, J Acosta-Uribe, A Baena, K J Wojta, G Coppola, R Fitch, M D de Both, M J Huentelman, E M Reiman, M E Brunkow, G Glusman, J C Roach, A W Kao, F Lopera & K S Kosik

doi:10.1038/mp.2015.131

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## Olfactory stem cells reveal MOCOS as a new player in autism spectrum disorders **OPEN**

F Féron, B Gepner, E Lacassagne, D Stephan, B Mesnage, M-P Blanchard, N Boulanger, C Tardif, A Devèze, S Rousseau, K Suzuki, J C Izpisua Belmonte, M Khrestchatisky, E Nivet & M Erard-Garcia

doi:10.1038/mp.2015.106

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## Direct conversion of human fibroblasts to induced serotonergic neurons **OPEN**

Z Xu, H Jiang, P Zhong, Z Yan, S Chen & J Feng

doi:10.1038/mp.2015.101

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## A genome-wide analysis of putative functional and exonic variation associated with extremely high intelligence **OPEN**

S L Spain, I Pedroso, N Kadeva, M B Miller, W G Iacono, M McGue, E Stergiakouli, G D Smith, M Putallaz, D Lubinski, E L Meaburn, R Plomin & M A Simpson

doi:10.1038/mp.2015.108

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## CNS-specific regulatory elements in brain-derived HIV-1 strains affect responses to latency-reversing agents with implications for cure strategies **OPEN**

L R Gray, D Cowley, C Welsh, H K Lu, B J Brew, S R Lewin, S L Wesselingh, P R Gorry & M J Churchill

doi:10.1038/mp.2015.111

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