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De novo heterozygous mutations in SMC3 cause a range of Cornelia de Lange syndrome-overlapping phenotype

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#	Paper	IF	Citations
64	Cohesin and human disease: lessons from mouse models. <i>Current Opinion in Cell Biology</i> , 2015 , 37, 9-17	9	27
63	Response to Dylan Mordaunt and Alisha McLauchlan. <i>Clinical Genetics</i> , 2015 , 88, 99-100	4	
62	Pathway Analysis Based on a Genome-Wide Association Study of Polycystic Ovary Syndrome. <i>PLoS ONE</i> , 2015 , 10, e0136609	3.7	27
61	TAF1 Variants Are Associated with Dysmorphic Features, Intellectual Disability, and Neurological Manifestations. <i>American Journal of Human Genetics</i> , 2015 , 97, 922-32	11	70
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59	Deletion of 11q12.3-11q13.1 in a patient with intellectual disability and childhood facial features resembling Cornelia de Lange syndrome. <i>Gene</i> , 2015 , 572, 130-134	3.8	8
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57	Biochemical and structural characterization of HDAC8 mutants associated with Cornelia de Lange syndrome spectrum disorders. <i>Biochemistry</i> , 2015 , 54, 6501-13	3.2	32
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37 36 35 34 33	Cohesin complex-associated holoprosencephaly. <i>Brain</i> , 2019 , 142, 2631-2643 A De novo HDAC2 variant in a patient with features consistent with Cornelia de Lange syndrome phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2019 , 179, 852-856 Next generation sequencing identified two novel mutations in NIPBL and a frame shift mutation in CREBBP in three Chinese children. <i>Orphanet Journal of Rare Diseases</i> , 2019 , 14, 45 Cornelia de Lange syndrome in diverse populations. <i>American Journal of Medical Genetics, Part A</i> , 2019 , 179, 150-158 Abnormal Body Size and Proportion. 2019 , 81-143 Another case of holoprosencephaly associated with RAD21 loss-of-function variant. <i>Brain</i> , 2020 ,	11.2 2.5 4.2 2.5	31 10 6 25

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