



Melanie M Y Chan

Contact

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Publications (1)

Mingardo E, Beaman G, Grote P, Nordenskjöld A, Newman W, Woolf A, Eckstein M, Hilger A, Dworschak G, Rösch W, Ebert A, Stein R, Brusco A, Di Grazia M, Tamer A, Torres F, Hernandez J, Erben P, Maj C, Olmos J, Riancho J, Valero C, Hostettler I, Houlden H, Werring D, Schumacher J, Gehlen J, Giel A, Buerfent B, Arkani S, Åkesson E, Rotstein E, Ludwig M, Holmdahl G, Giorgio E, Berettini A, Keene D, Cervellione R, Younsi N, Ortlieb M, Oswald J, Haid B, Promm M, Neissner C, Hirsch K, Stehr M, Schäfer F, Schmiedeke E, Boemers T, van Rooij I, Feitz W, Marcelis C, Lacher M, Nelson J, Ure B, Fortmann C, Gale D, Chan M, Ludwig K, Nöthen M, Heilmann S, Zwink N, Jenetzky E, Odermatt B, Knapp M, Reutter H. A genome-wide association study with tissue transcriptomics identifies genetic drivers for classic bladder exstrophy. *Commun Biol* 2022; 5:1203.

Projects (0)

No results found.

Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

support.forschung@kssg.ch