Title: *STXBP1* Encephalopathy with Epilepsy *GeneReview* – Phenotypes Associated with Recurrent Pathogenic Variants in *STXBP1* 

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## Phenotypes Associated with Recurrent Pathogenic Variants in STXBP1

Some *STXBP1* pathogenic missense variants occurring in more than one affected individual were listed below, comparing phenotypes:

- c.1217G>A was found in eight unrelated individuals. All had severe to profound global developmental delay or intellectual disability (ID). Seizures required two or more anti-epileptic drugs (AEDs) or were drug resistant [Saitsu et al 2010, Mignot et al 2011, Allen et al 2013, Di Meglio et al 2015, Romaniello et al 2015, Stamberger et al 2016].
- c.875G>A was found in five unrelated individuals [Michaud et al 2014, Helbig et al 2016, Stamberger et al 2016, Trump et al 2016]. Age at seizure onset varied from 6 months to 8 years. Seizure types included infantile spasms and partial seizures. ID ranged from mild to severe. Two individuals had a good response to AEDs becoming seizure free [Stamberger et al 2016].
- c.703C>T was found in three unrelated individuals. All had severe to profound global developmental delay or ID. Two became seizure free [Saitsu et al 2010, Allen et al 2013, Boutry-Kryza et al 2015].
- c.364C>T was found in four unrelated individuals [Lemke et al 2012, Mercimek-Mahmutoglu et al 2015, Stamberger et al 2016]. Age of seizure onset varied between the neonatal period and age one year. One individual had no seizure history.
- c.1439C>T was found in three unrelated individuals. All had severe
  developmental delay. Two became seizure free and one was refractory to antiseizure treatment [Milh et al 2011, Tso et al 2014, Di Meglio et al 2015].
- c.703C>T was found in three unrelated individuals. All had severe to profound global developmental delay or ID. Two became seizure free [Saitsu et al 2010, Allen et al 2013, Boutry-Kryza et al 2015].
- c.568C>T, c.416C>T, c.1162C>T, c.1651C>T, and c.902+1G>A were each found in two or three unrelated individuals for whom no detailed phenotypic information was available [Hamdan et al 2009, Milh et al 2011, Kodera et al 2013, Allen et al 2013, Weckhuysen et al 2013, Barcia et al 2014, Michaud et al 2014, Di Meglio et al 2015, Keogh et al 2015, Helbig et al 2016, Li et al 2016].

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