

## Supplemental Data

### Mutations in *GBA2* Cause Autosomal-Recessive

### Cerebellar Ataxia with Spasticity

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**Figure S1.** Output of homozygosity mapper for families C, E and I across the whole genome. Data input comprised genotypes from the HumanOmniExpress12 array (~730,000 markers/sample). Red lines indicate contiguous stretches of homozygous genotypes identical by descent in the 7 affected individuals.

