



Figure 1. Worldwide distribution of Spinocerebellar Ataxia (SCA) subtypes [Schöls et al 1997, Moseley et al 1998, Saleem et al 2000, Storey et al 2000, Tang et al 2000, Maruyama et al 2002, Silveira et al 2002, van de Warrenburg et al 2002, Dryer et al 2003, Brusco et al 2004, Schöls et al 2004, Shimizu et al 2004, Zortea et al 2004, Jiang et al 2005, Jiang et al 2013]

Note: The data in Figure 1 was gathered before the use of multigene panels and exome/genome sequencing. The data are still appropriate for nucleotide repeat disorders; new SCAs found by exome/genome sequencing represent a very small percentage. There have been no recent regional updates to SCA subtype distribution in Australia or sub-Saharan Africa.