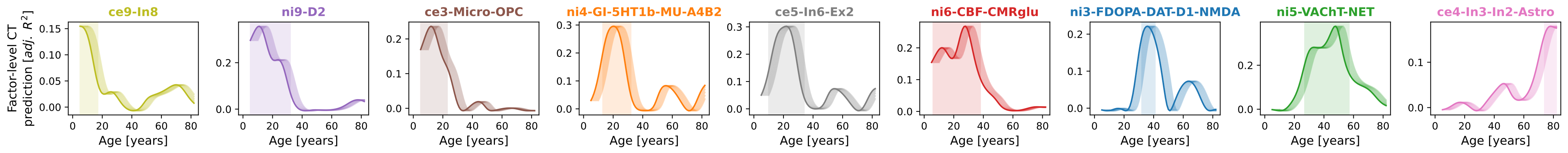


CT change explained by factor-level neurobiological markers



Gene expression associated to original atlases that load on each factor

