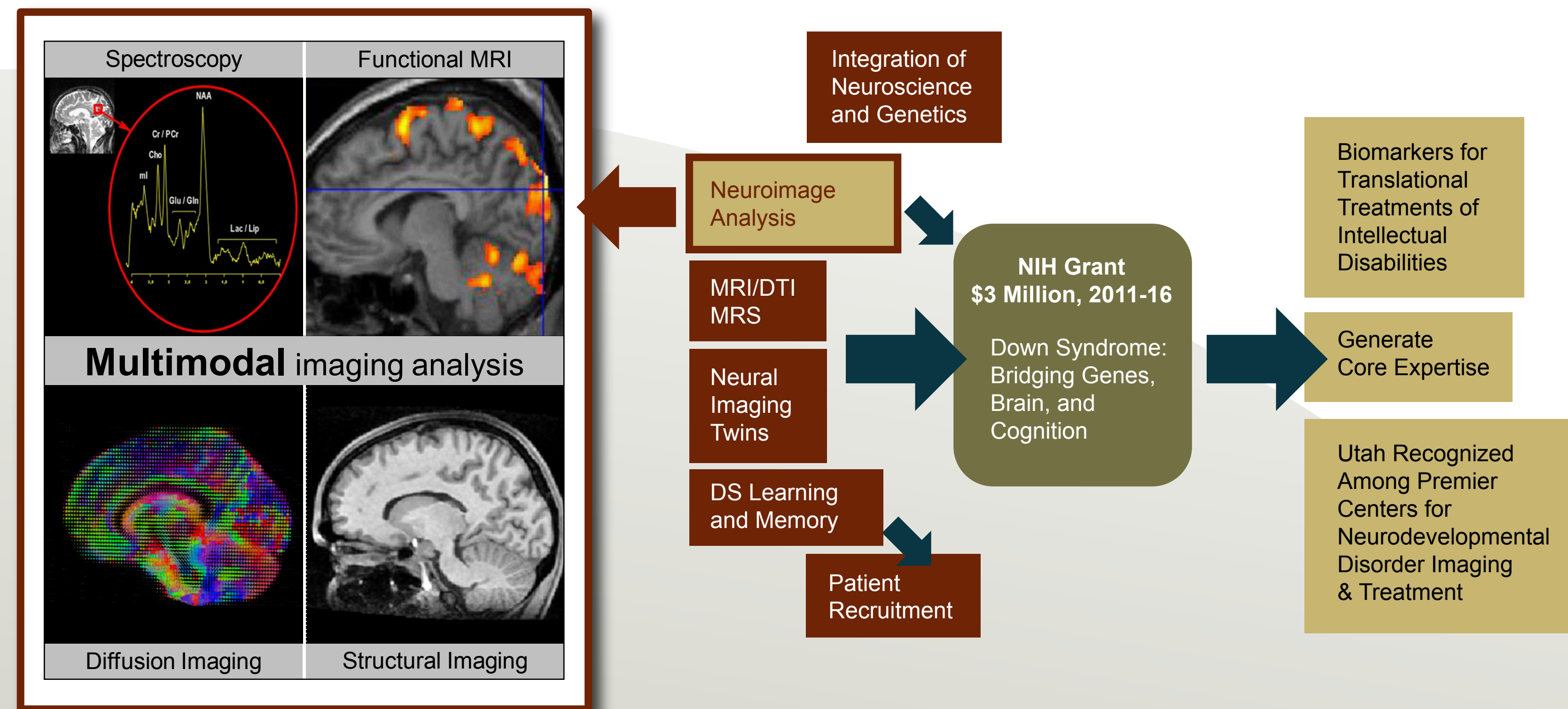
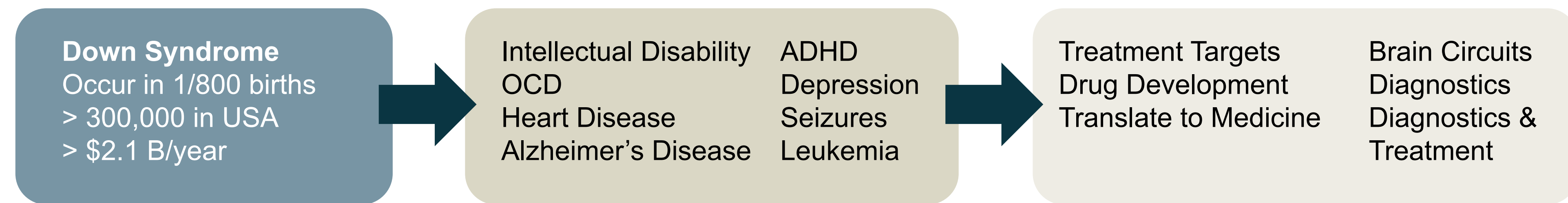


Down syndrome: Bridging Genes, Brain and Cognition

Sylvain Gouttard, Julie Korenberg, Guido Gerig

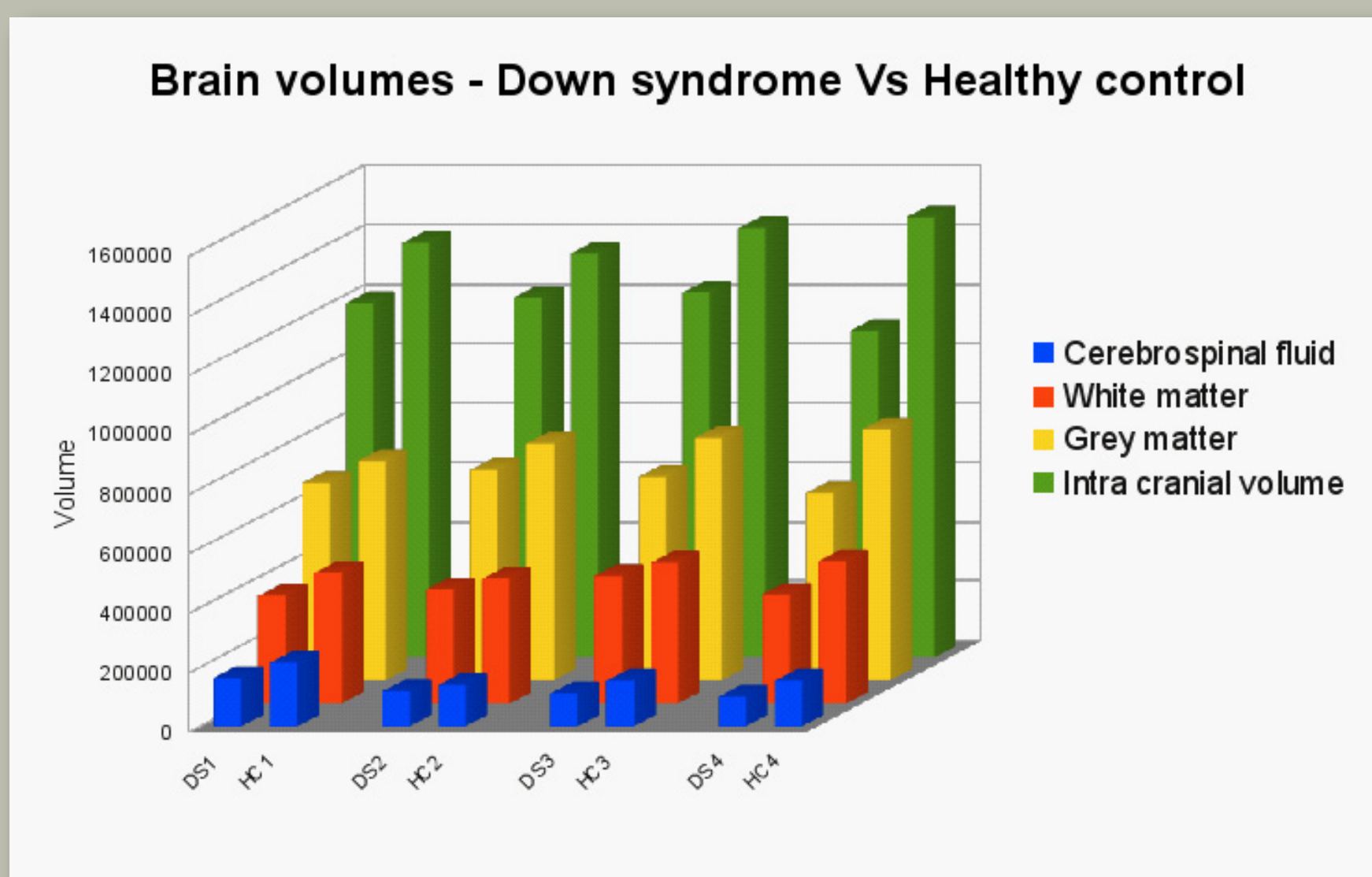


Brain volume analysis

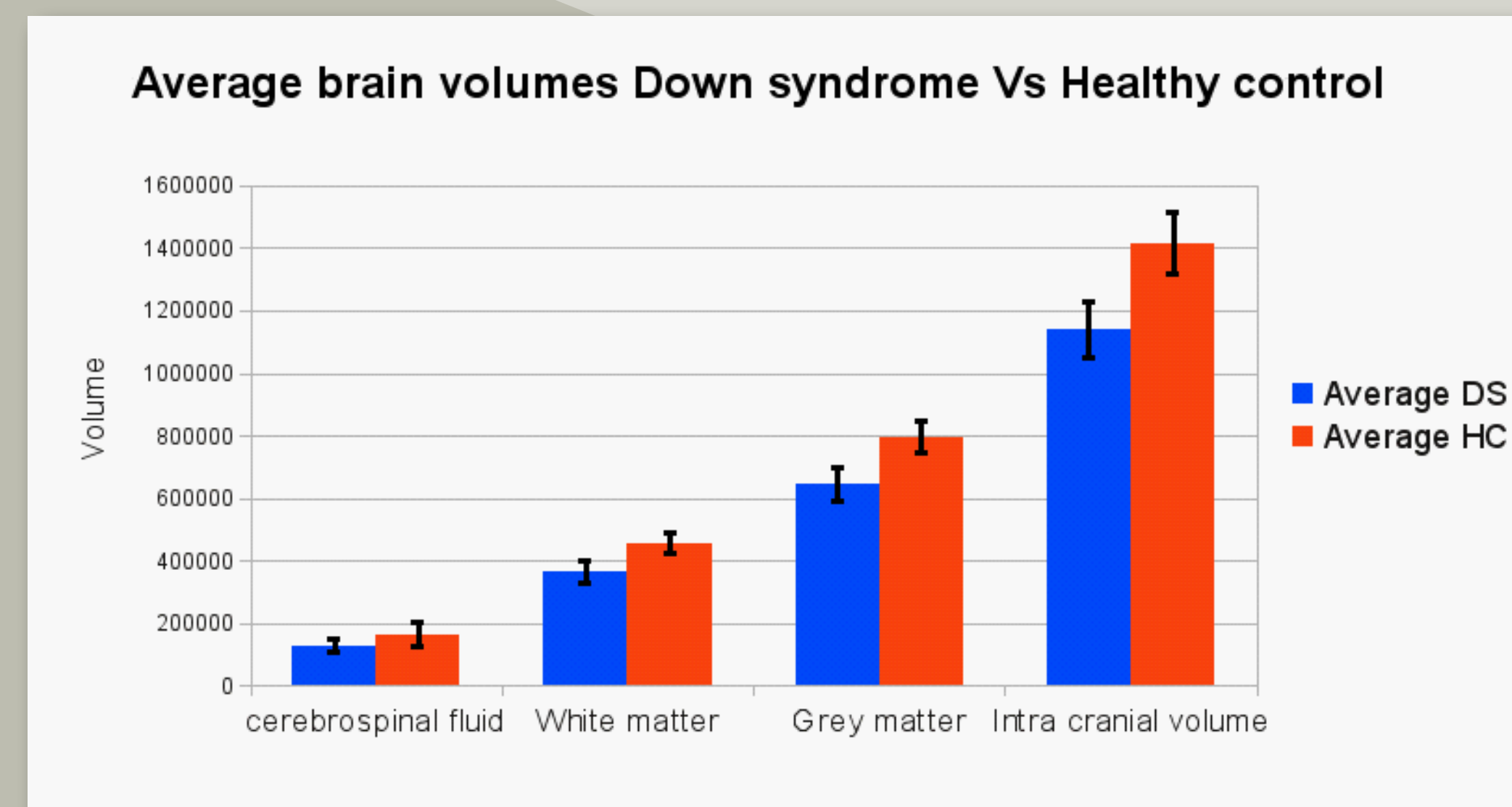
Typical Down Syndrome phenotype includes different trajectory of neuroanatomy development. Analysis of these differences will give insight into genotype/phenotype/behavior relationships and influence of genetics on brain development.

Analysis performed on 8 Down syndromes (DS) and 9 healthy subjects (HC).

- ▶ Average brain volumes of the 2 population shows significant differences for all tissues.
- ▶ Age/sex paired subjects (DS vs HC) show consistent differences.



Volumes of brain tissues, fluid and intra-cranial space for four pairs of age and sex matched subjects: one Down syndrome and one healthy subject



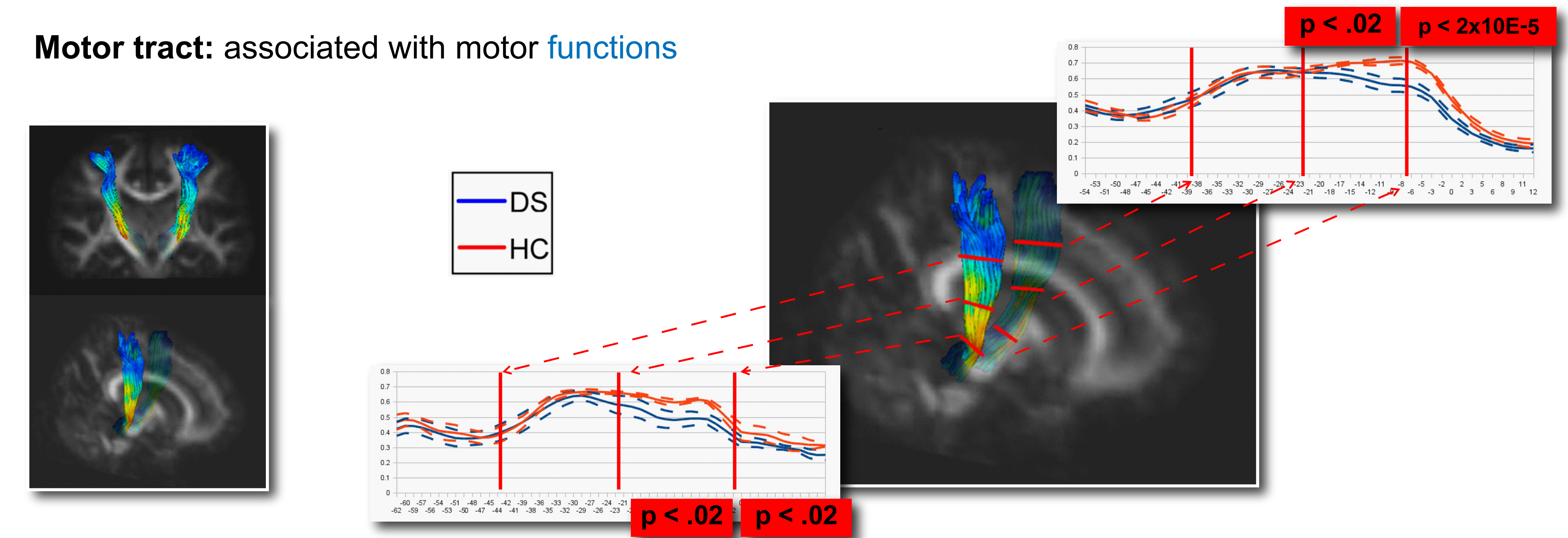
Average volumes of brain tissues, fluid and intra-cranial space for the 2 populations (8 DS and 9 HC).

White matter integrity analysis

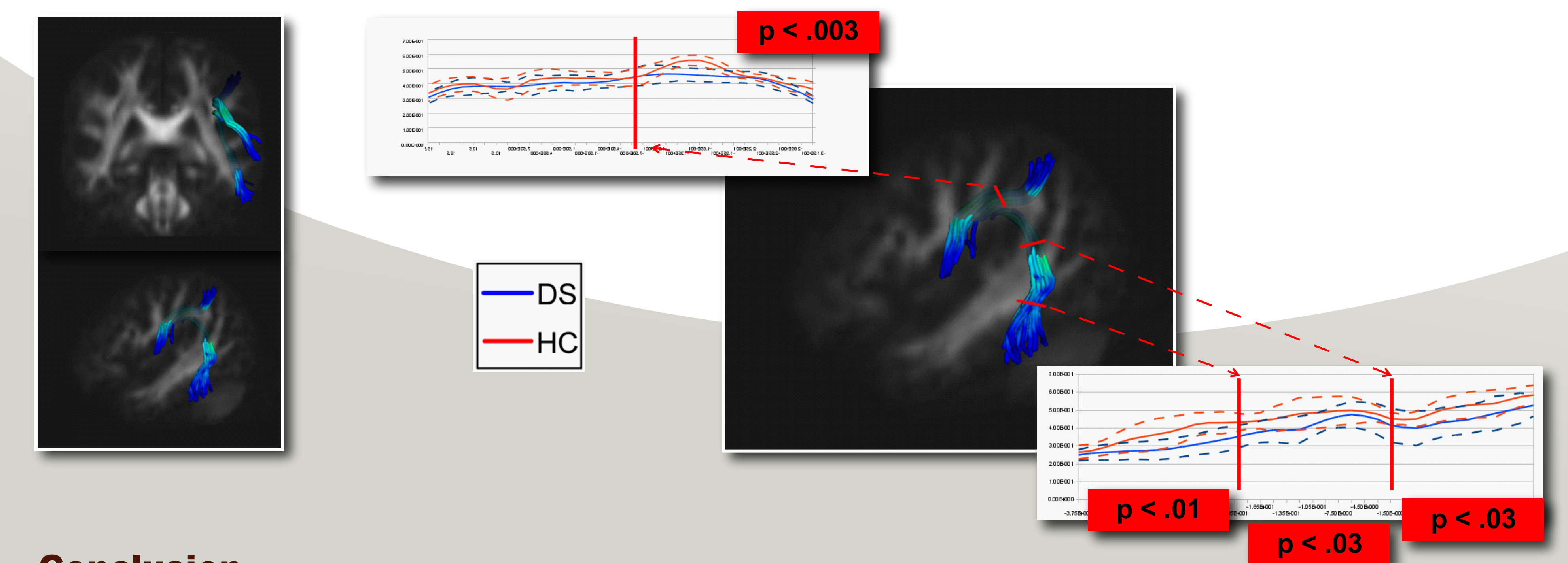
Down syndrome subjects show **cognitive differences** that can be characterized in the white matter integrity of specific brain connectivity pathways. Brain white matter analysis can highlight relationship of genotype and brain connectivity analysis.

- ▶ Most portions analyzed tracts show highly significant differences between groups.

Motor tract: associated with motor functions



Arcuate tract: associated with language



Conclusion

Volumetric analysis demonstrates **significant differences of the trajectories** of brain anatomy development in DS subjects → Refined exploration of localized volume and shape differences with help to define new biomarkers of disease.

White matter integrity analysis **reflects the observed cognitive differences** between Down syndrome and control groups → White matter tract analysis might serve as a biomarker of specific aspects of cognitive development.

