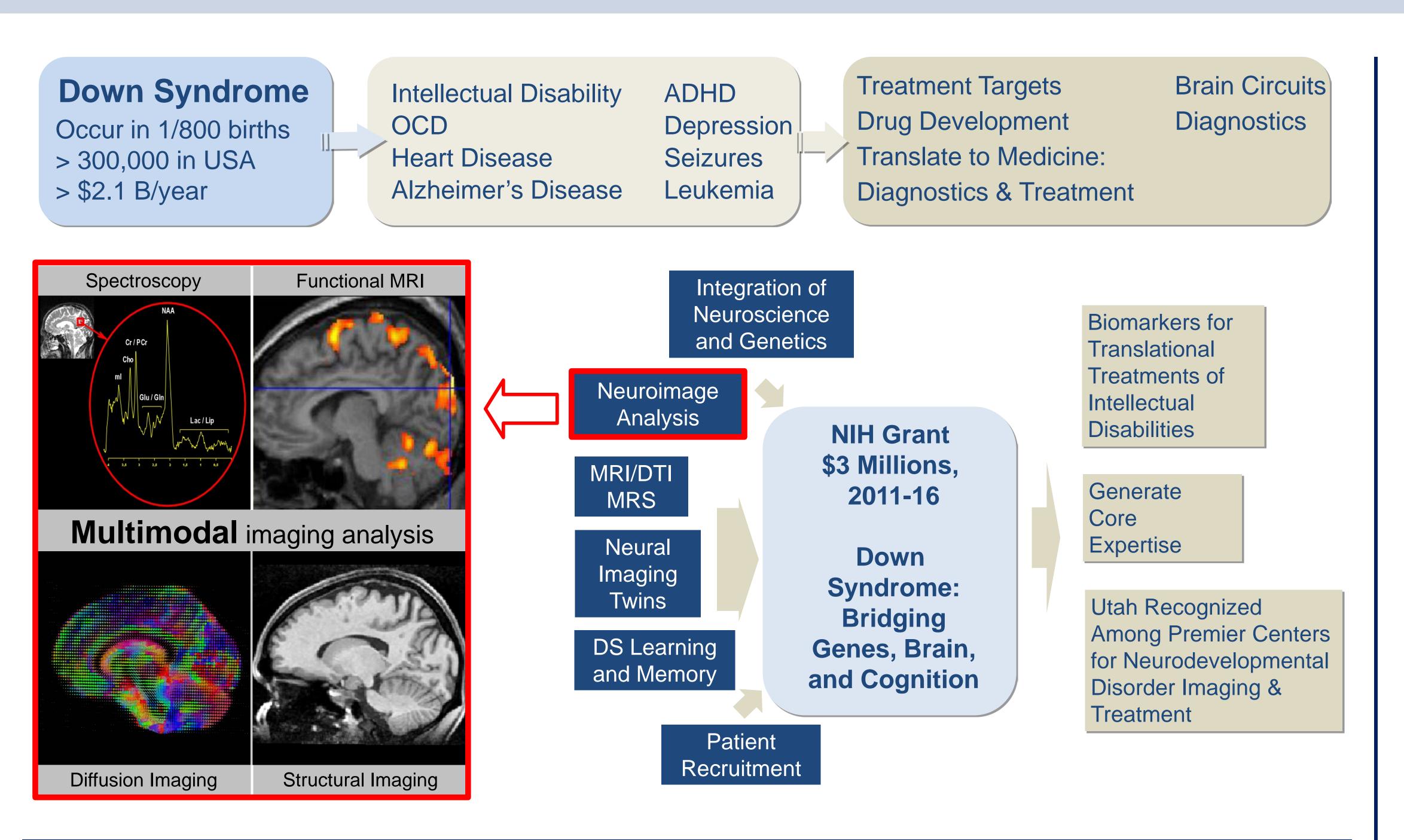
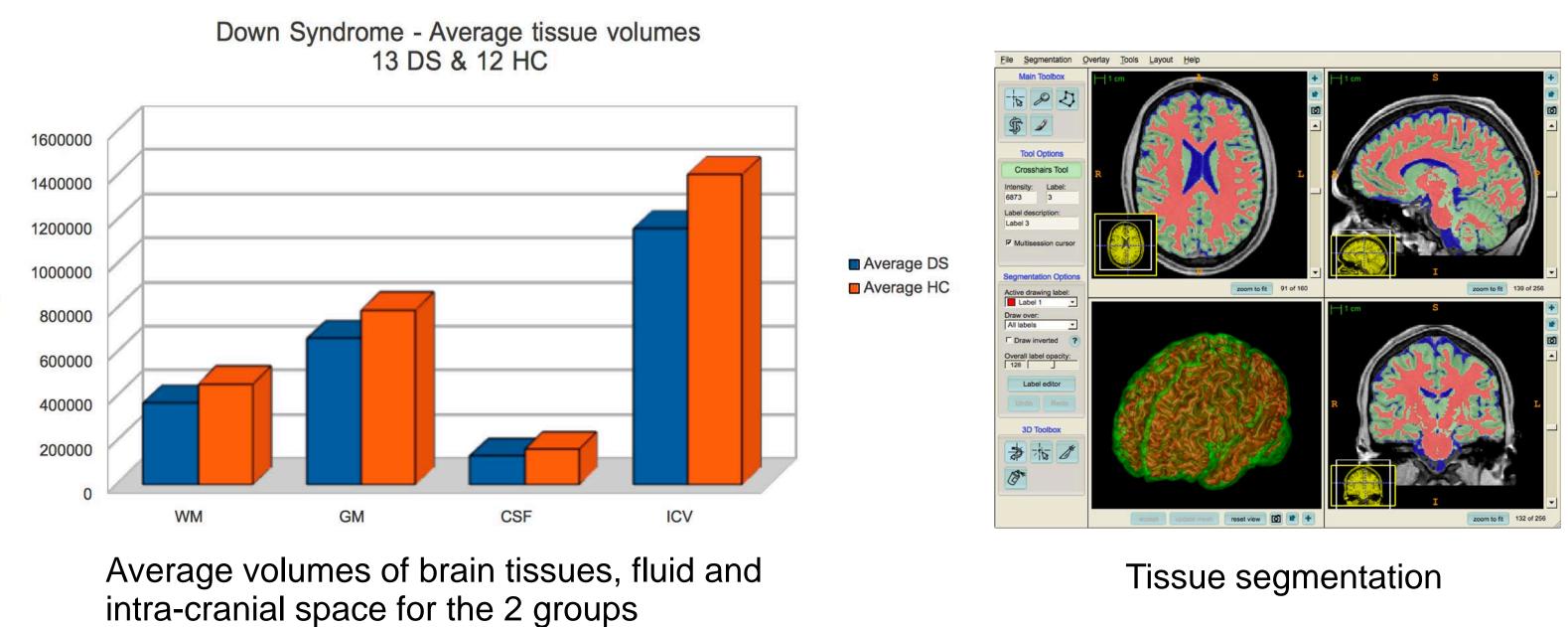
Clement Vachet, Sylvain Gouttard, Julie Korenberg, Guido Gerig



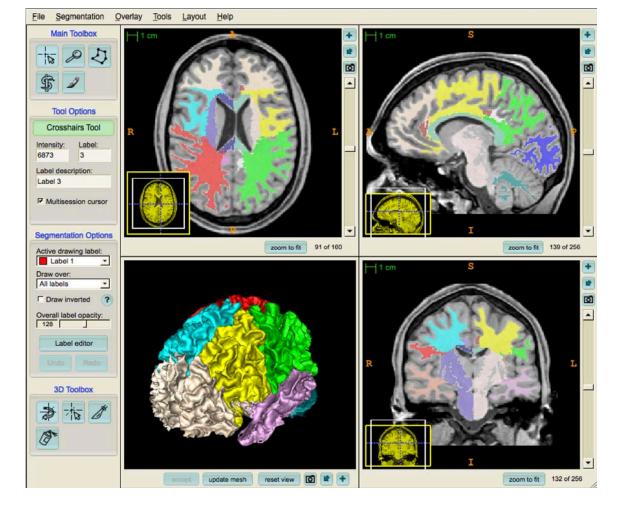
Brain structural analysis

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

Analysis performed on 13 Down syndromes (DS) and 12 healthy subjects (HC): => Average lobar tissue volumes show significant group differences. => Average subcortical structures volumes show significant group differences.



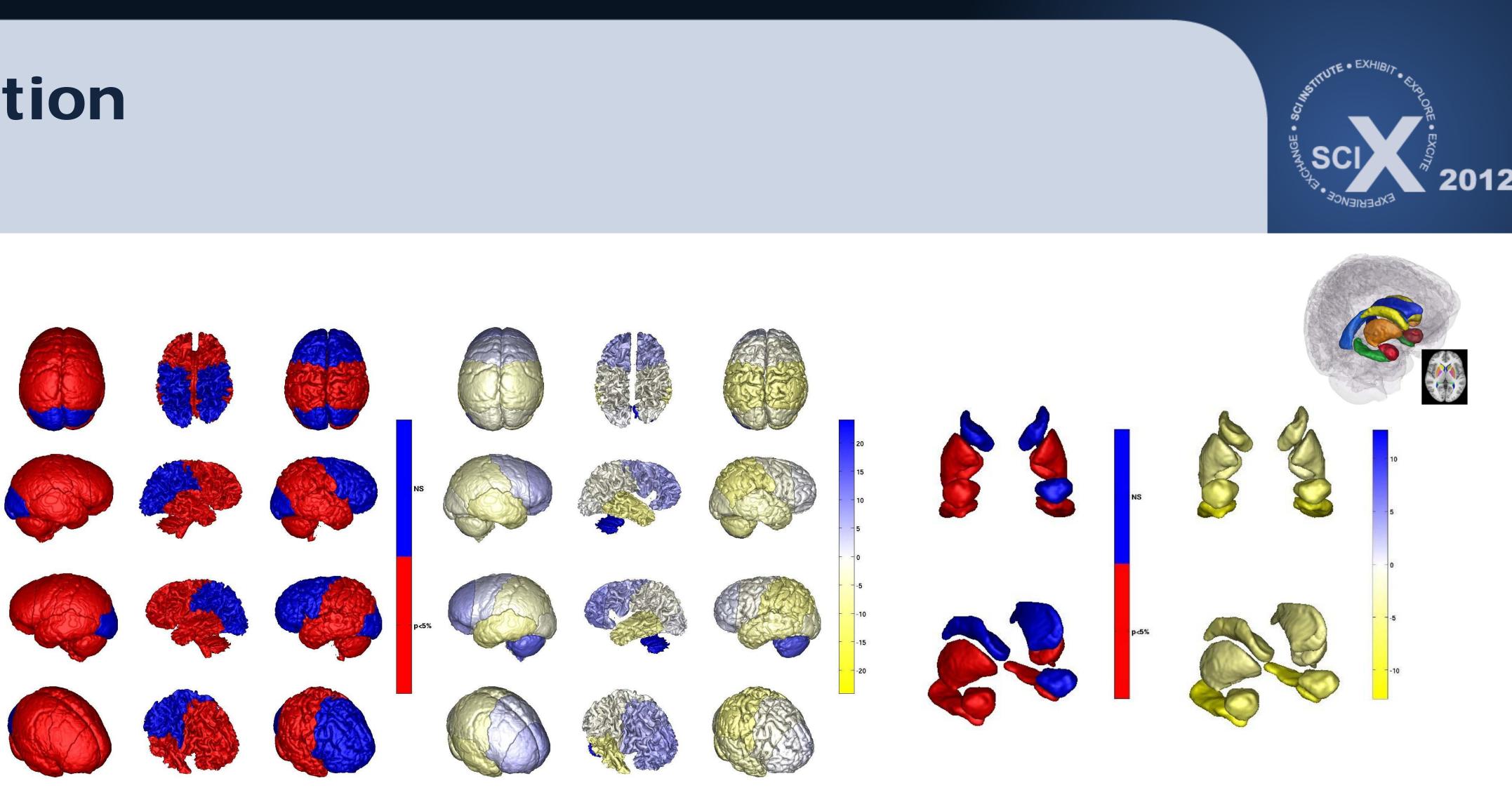




Lobar white matter segmentation





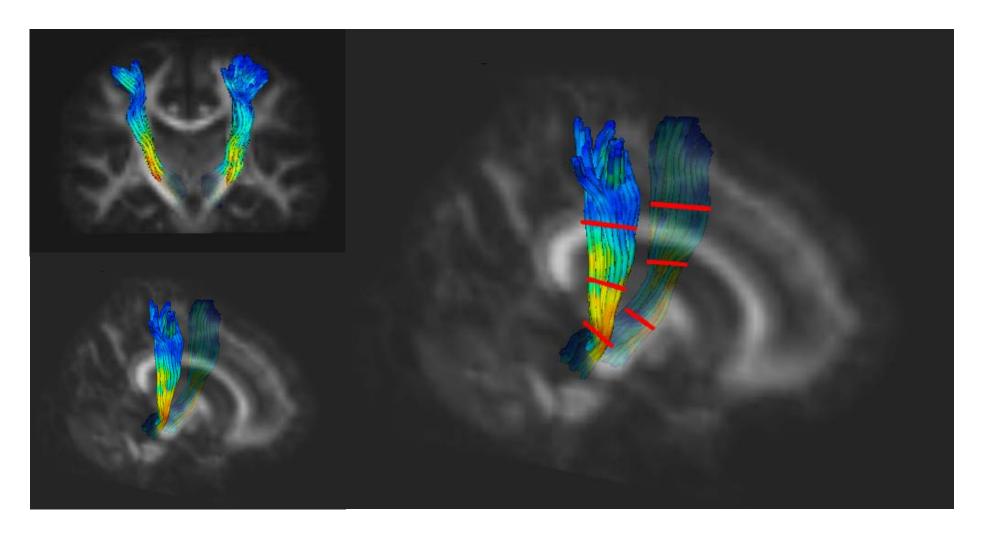


Significance maps (LEFT) and ICV normalized lobar volume changes in percent between healthy controls (HC) and DS (RIGHT: blue: HC > DS, yellow: HC < DS)

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 of analyzed tracts show highly significant group differences (lower fractional anisotropy in DS group).

Motor tract: associated with motor functions



Conclusion

Structural volumetric analysis demonstrates significant differences of the **trajectories** of brain anatomy development in DS subjects => Refined exploration of localized volume and shape differences will help 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.





Significance maps (LEFT) and ICV normalized subcortical volume changes in percent between controls and DS (RIGHT: yellow: HC < DS)

Arcuate tract: associated with language

