

# Documentation

## Documentation: Allen Mouse Brain Atlas

Document	Description
<a href="#">In Situ Hybridization</a>	Processes and procedures used to perform ISH
<a href="#">Informatics Data Processing</a>	Design and implementation of the Informatics Data Pipeline (IDP)
<a href="#">Reference Atlas, Version 1 (2008)</a>	Overview of the sagittal and coronal Allen Reference Atlases
<a href="#">Reference Atlas, Version 2 (2011)</a>	Overview of the sagittal and coronal Allen Reference Atlases
<a href="#">Mouse CCF, Reference Atlas, Version 3 (2015)</a>	Overview of the design and implementation of the Allen Mouse Common Coordinate Framework
<a href="#">Fine Structure Annotation</a>	Explanation of the informatics methods used to create the fine structure annotations
<a href="#">Acknowledgement of Collaborators</a>	Recognition of collaborators and others who provided assistance

## Supplemental Data: Allen Mouse Brain Atlas

Document	Description
<a href="#">ISH Platform Controls</a>	Quality control standards for high-throughput RNA in situ hybridization data generation
<a href="#">Cross Platform Validation</a>	Methods for validating the data
<a href="#">AGEA User Guide</a>	Overview of the Anatomic Gene Expression Atlas (AGEA)
<a href="#">Sleep Study</a>	PDF Article of the Sleep Study Results
<a href="#">Mouse Strains Study</a>	Mouse Strains Study
<a href="#">FSA Reports</a>	Compilations of neuroanatomic and gene expression characteristics for key brain regions or nuclei
<a href="#">Somatosensory Annotation</a>	Technical white paper detailing the methods used to manually annotate the expression patterns of genes in the mouse somatosensory cortex
<a href="#">Comparison of the Top 1000 Genes</a>	The 1000 most requested genes were analyzed as a validation of the ISH data
<a href="#">Comparison of the Top 1000 Genes (Oct 2010)</a>	An updated list of the 1000 most requested genes that were analyzed as a validation of the ISH data