

Finding expressed genes and novel phyto-compounds as ligands from schizophrenic convergent genomic and pharmacological evidence of PI3K/GSK3 signaling alterations in neurons

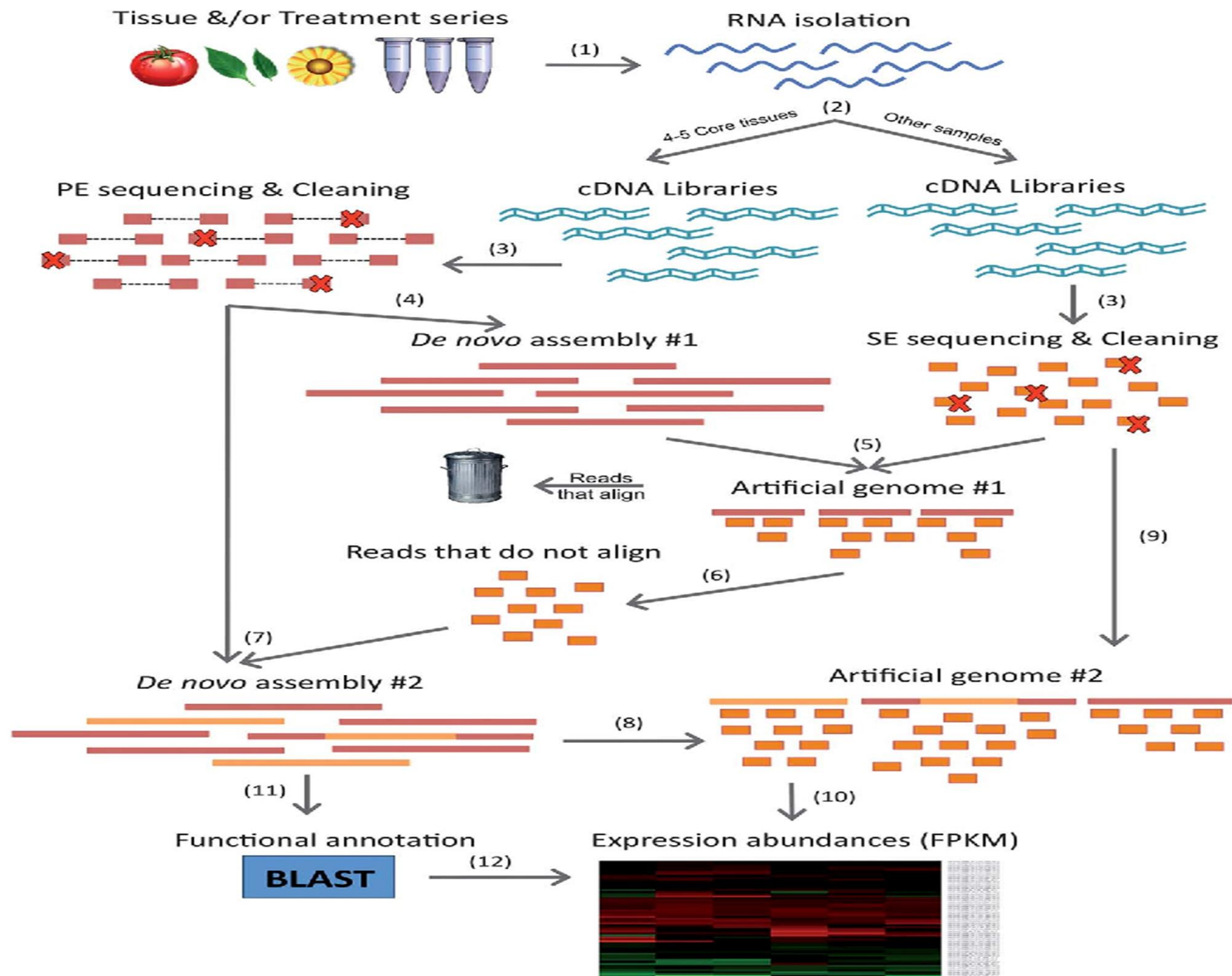
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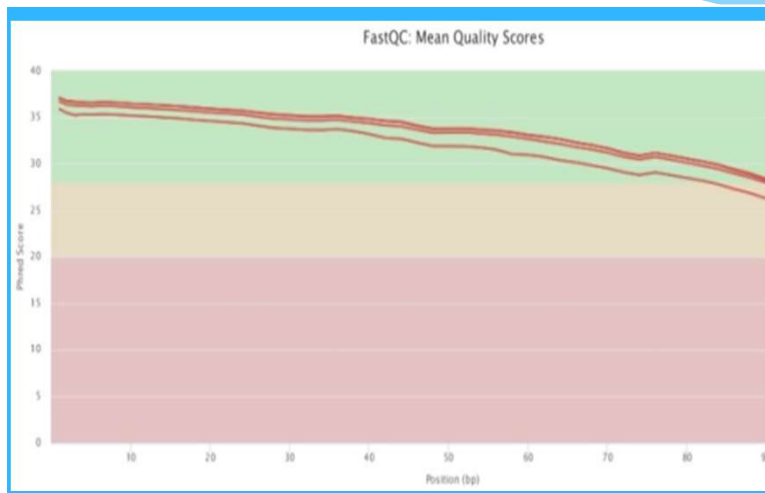
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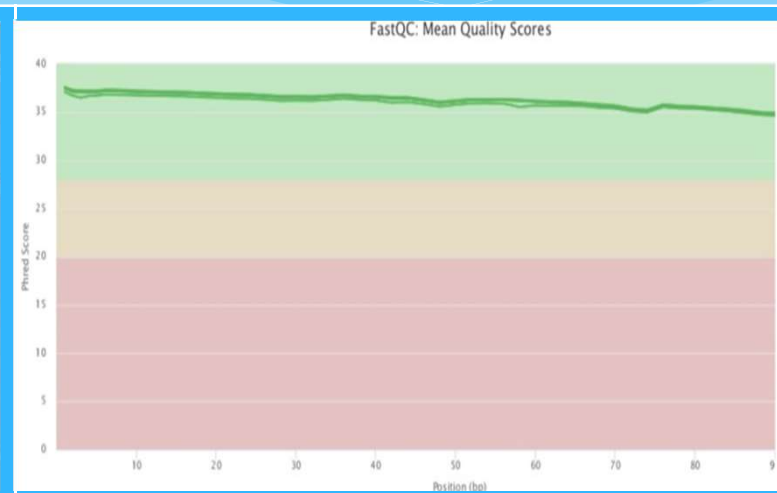
Data retrieval

- * Schizophrenia RNA-seq data SRR15629349, SRR15629350, SRR15629351 and SRR15629352 were retrieved from SRA database.
- * Work by Stertz L *et al.*, "Convergent genomic and pharmacological evidence of PI3K/GSK3 signaling alterations in neurons from schizophrenia patients.", *Neuropsychopharmacology*, 2021 Feb;46(3):673-682

Quality checking of the sequences

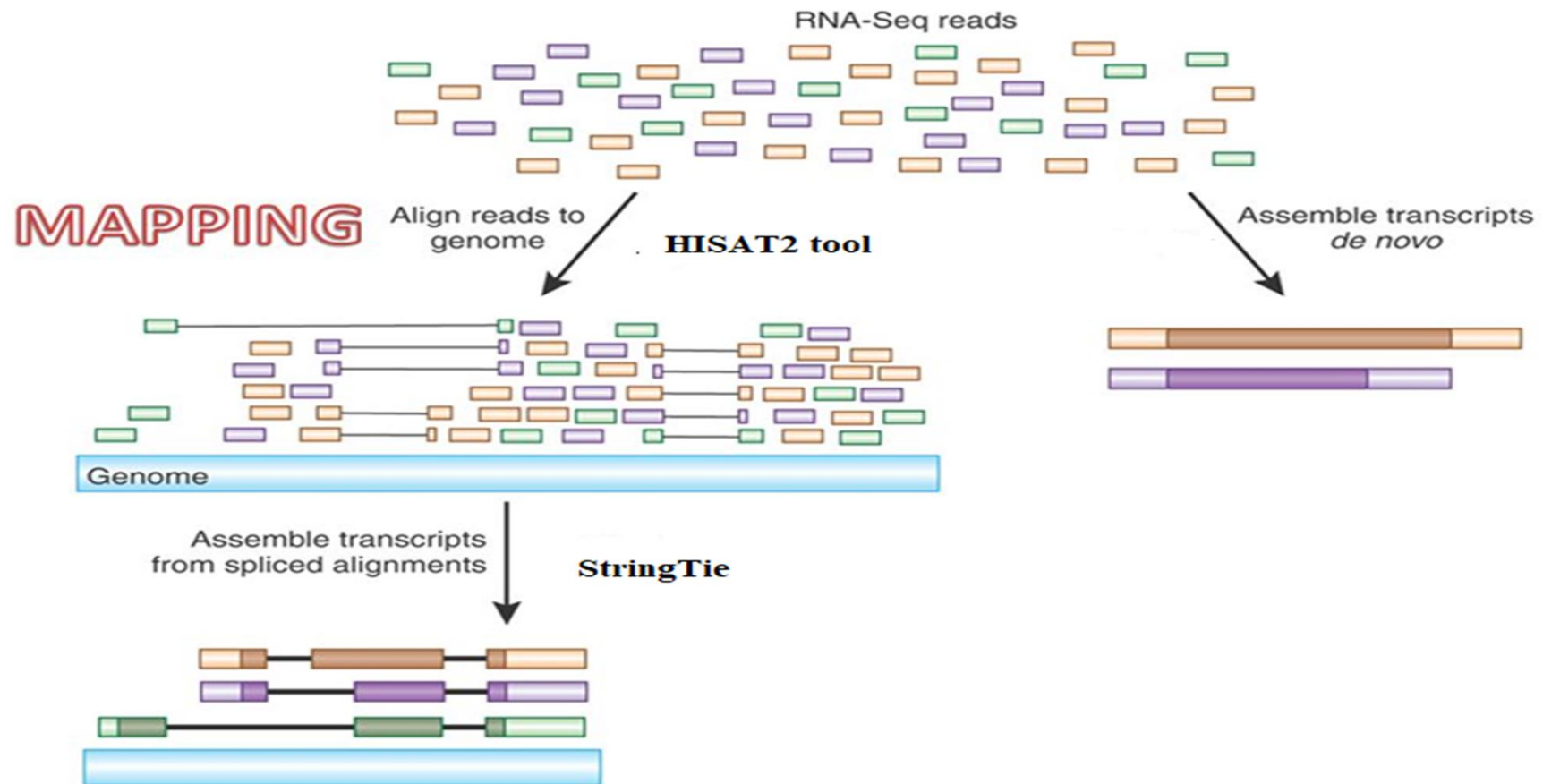


Raw sequence quality

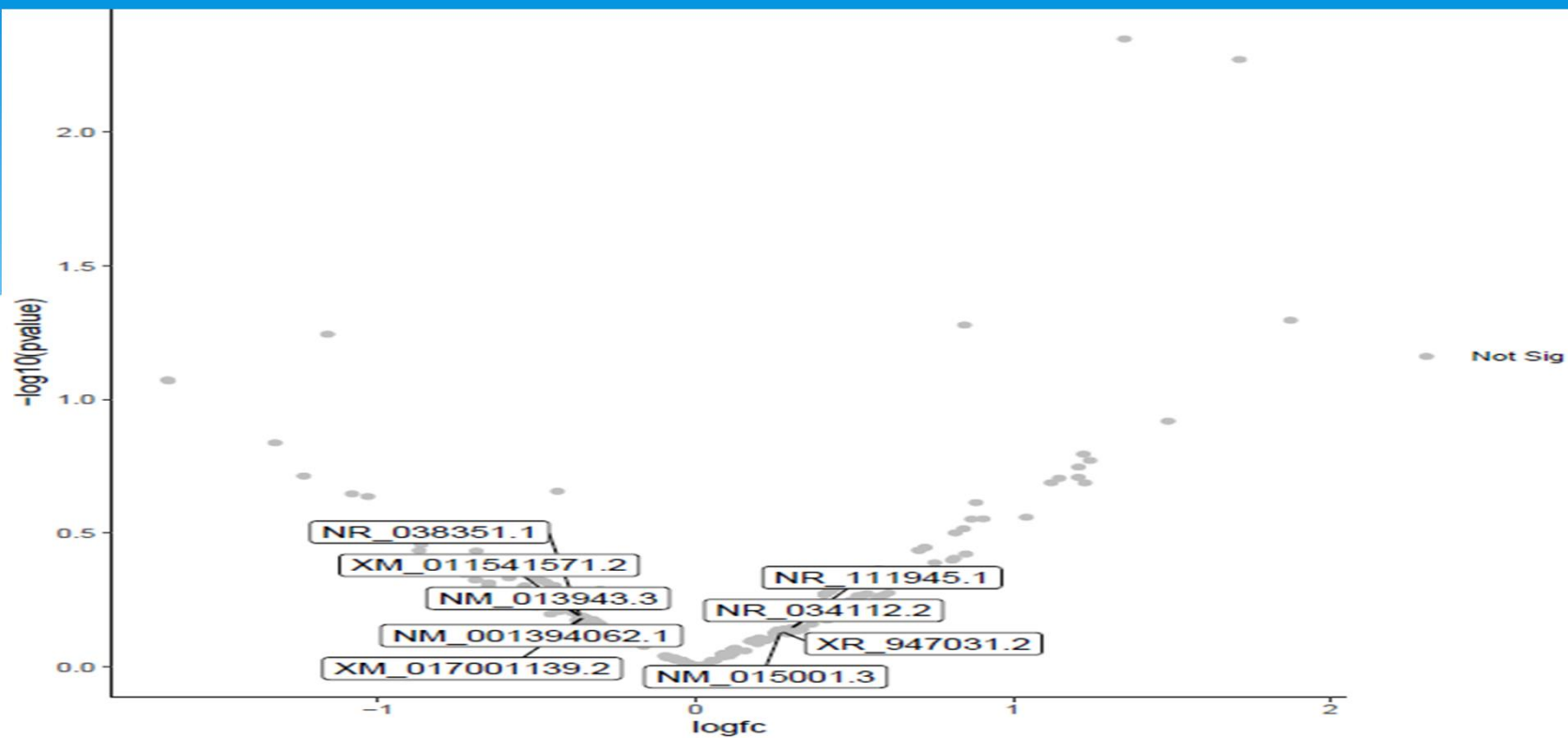


Sequence quality after trimming

Sequence quality was checked before and after trimming. This was done by tool FASTQC.



- ❖ The sequences were mapped with human reference genome using HISAT2 tool
- ❖ Transcriptomes were generated with Stringtie
- ❖ Transcriptome database was made by using tool Stringtie - Merge
- ❖ Next, GFFcompare annotated the transcripts of our newly created transcriptome to make us known the relationship of each transcript to the human RefSeq reference
- ❖ Further, using FeatureCounts tool the reads aligning in exons of our GFFCompare generated transcriptome database was counted



- DESeq tool was used to identify the expressed genes.
- NR_111945.1, NR_034112.2, XR_947031.2 and NM_015001.3 which are towards the right of the volcano plot are the most up-regulated genes and NM_001039211.3, NR_038351.1, XM_011541571.2, NM_013943.3, XM_017001139.2 and NM_001394062.1 are towards the left are the most down-regulated genes
- From the genes above, NR_038351.1 is the most significant gene since it is on the topmost amongst the genes.

High-Throughput Screening (HTS)

- * Homology modeling of the selected above receptors was done.
- * Phyto-compounds from Ayurvedic Medicinal plants *Withania somnifera*, *Melisa officinalis*, *Piper methysticum*, *Valeriana officinalis* and *Hypericum perforatum* are traditionally used to treat many diseases are considered here.
- * As per Lipinski's rule of five [ADME (Adsorption, distribution and metabolism extraction)] we checked the drug likeliness of the above phyto-compounds.
- * As per docking result it is seen that the phytochemicals Lutolin, Kaempferol, Ginkgolide A, Ascorbic acid, Tetrahydroxyangonin, Matricarinol, 5-hydroxy-4,7-dimethoxy, Tetrahydroxyangonin, D-Allose and Azetidine docks with all the receptors.
- * Hence these compounds can be successfully considered as ligands for the receptors on further validation using in-vitro and in-vivo studies.

Reference

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