| Sl.No. | Transcriptome   | SRA/GEO ID     | Plant                |
|--------|---|----------------|----------------------|
| 1      | Histological and microarray analysis of the direct effect of water shortage alone or combined with heat on early grain development in wheat ( <i>Triticum aestivum</i> ). Physiologia Plantarum 2012; 140: 174–188.   | GSE18205       | Wheat                |
| 2      | Transcriptomic analysis of Sorghum bicolor responding to combined heat and drought stress. BMC Genomics 2014; 15:456.   | GSE48205       | Sorghum              |
| 3      | Unique Physiological and Transcriptional Shifts under Combinations of Salinity, Drought and Heat. Plant physiology 2017; DOI:10.1104/pp.17.00030.   | PRJNA360513    | Arabidopsis          |
| 4      | Water deficit modulates the response of <i>Vitis vinifera</i> to the Pierce's disease pathogen <i>Xylella fastidiosa</i> . Mol Plant-Microbe Interactions 2013;26: 643–657.   | GSE44213       | Vitis vinifera       |
| 5      | Identification of genes involved in the response of Arabidopsis to simultaneous biotic and abiotic stresses. Plant Physiology 2013; 162: 2028–2041.   | NASCARRAYS-489 | Arabidopsis          |
| 6      | Combined biotic stresses trigger similar transcriptomic responses but contrasting resistance against a chewing herbivore in Brassica nigra  | E-MTAB-5030    | Brassica nigra       |
| 7      | Transcriptome dynamics of Arabidopsis during sequential biotic and abiotic stresses   | PRJNA315516    | Arabidopsis          |
| 8      | Pre-exposure of Arabidopsis to the abiotic or biotic environmental stimuli "chilling" or "insect eggs" exhibits different transcriptomic responses to herbivory   | GEO- GPL19779  | Arabidopsis          |
| 9      | Effect of prior drought and pathogen stress on Arabidopsis transcriptome changes to caterpillar herbivory   |                | Arabidopsis          |
| 10     | A step towards understanding plant responses to multiple environmental stresses: a genome-wide study  | GSE39956       | Arabidopsis          |
| 11     | Transcriptome Responses to Combinations of Stresses in Arabidopsis  | GSE41935       | Arabidopsis          |
| 12     | SKM107: Drought and nonhost pathogen  |                | Arabidopsis          |
| 13     | SKM107: Drought and host pathogen   |                | Arabidopsis          |
| 14     | Dual impact of elevated temperature on plant defence and bacterial virulence in Arabidopsis   | PRJNA325245    | Arabidopsis          |
| 15     | SKM107: Host pathogen & drought   |                | Arabidopsis          |
| 16     | SKM107: Drought recovery & pathogen   |                | Arabidopsis          |
| 17     | Transcriptomic reprogramming of barley seminal roots by combined water deficit and salt stress  | SRP133479      | Barley               |
| 18     | Physiological analysis and transcriptome sequencing reveal the effects of combined cold and drought on tomato leaf  | SRP156535      | Tomato               |
| 19     | Zheng C, Wang Y, Ding Z and Zhao L (2016) Global<br>Transcriptional Analysis Reveals the Complex<br>Relationship between Tea Quality, Leaf Senescence and<br>the Responses to Cold-Drought Combined Stress in<br>Camellia sinensis. Front. Plant Sci. 7:1858. | SRP091321      | Camellia<br>sinensis |

| 20 | Kumar, D., Hazra, S., Datta, R., & Chattopadhyay, S. (2016). Transcriptome analysis of Arabidopsis mutants suggests a crosstalk between ABA, ethylene and GSH against combined cold and osmotic stress. Scientific reports, 6, 36867              | GSE77490             | Arabidopsis                   |
|----|---|----------------------|-------------------------------|
| 21 | Kumar, D., Datta, R., Hazra, S., Sultana, A.,<br>Mukhopadhyay, R., & Chattopadhyay, S. (2015).<br>Transcriptomic profiling of Arabidopsis thaliana mutant<br>pad2.1 in response to combined cold and osmotic<br>stress. PloS one, 10(3), e0122690 | GSE61170             | Arabidopsis                   |
| 22 | ABA Is Required for Plant Acclimation to a Combination of Salt and Heat Stress  | GSE72806             | Arabidopsis                   |
| 23 | Unique Physiological and Transcriptional Shifts under Combinations of Salinity, Drought and Heat. Plant physiology 2017; DOI:10.1104/pp.17.00030.   | NCBI-<br>PRJNA360513 | Brachypodium dystachion       |
| 24 | Elevated ozone alters soybean-virus interaction   |                      | Soybean                       |
| 25 | Rice pyramided line IRBB67 (Xa4/Xa7) homeostasis under combined stress of high temperature and bacterial blight   | GSE79011             | Rice pyramided                |
| 26 | SKM107: Drought and host pathogen   |                      | Chickpea                      |
| 27 | Deep Transcriptome Sequencing of Wild Halophyte Rice,<br>Porteresia coarctata, Provides Novel Insights into the<br>Salinity and Submergence Tolerance Factors   | GSE44913             | Rice, Porteresia<br>coarctata |
| 28 | Transcriptomic profiling of Solanum peruvianum LA3858 revealed a Mi-3-mediated hypersensitive response to Meloidogyne incognita   | PRJNA494774          | Solanum<br>peruvianum         |
| 29 | Drought and flooding have distinct effects on herbivore-induced responses and resistance in Solanum dulcamara   | GSE69648             | Solanum<br>dulcamara          |