Work Distribution List (Final Table)

Tauseeq	Ahmad Zahoor	Azhar Bilal	Maaz
1. bx-python	26. BB Tools	51. RUST	76. PROTEO FORMER
2. Inchworm	27.SMALT	52. RiboProfiling	77. bzip2
3. SCLUST	28. Lifescope	53. riboWaltz	78. GNU zip
4. LULU	29. Kronos	54. Xtail	79. GDC
5.XenofilteR	30. ScaleHD	55. diricore	80. Biostrings
6. SUSHI	31. aRNApipe	56. Anota2seq	81. Anduril
7. Goleft	32. STpipeline	57. RIVET	82. vcfsubsample
8. AfterQC	33.Potcullis	58. Riborex	83. FASTX-Toolkit
9. PBcR	34. Scavenger	59 RiboDiff	84. Maftools
10. SOAPec	35. Splicify	60. Rfoot	85. Pgltools
11. MiRCA	36. Stringtie	61. ROSE	86. UBU
12. Jabba	37. Agalma	62. Toil	87. VCFcomparator
13. GotCloud	38. Nextflow	63. Ribomap	88. GenePattern
14. systemPipeR	39. Horse_trans	64. RiboAsiteDeblur	89. ea-utils
15. RSCU RS	40. QuickNGS	65. riboShape	90. Seqtk
16. QuickNGS	41. HTseq	66. RiboPip	91. FASconCAT
17. Shoelaces	42. RSeqC	67 RiboRL	92. Pysam
18.Scrappie	43. Qualimap	68. RiboTaper	93. ChIP-Extract
19.DeepNano	44. MultiQC	69 riboHMM	94. affyio
20. Metrichor	45. Phyluce	70. SPECtre	95. htslib
21.TagDust	46. Cd-hit-454	71. RiboProfiling	96. Cdbfasta
22.seqyclean	47. Nextclip	72. Rp-Bp	97. bedr
23. SEPia	48. samblaster	73. uORF-Tools	98.GFF3sort
24. Seqpurge	49. biobambam	74. REPARATION	99. Seqtk
25. BAMstats	50.expHTS	75. RiboCode	100. fastq-tools