

Bangladesh Field RNA-Seq Raw Data July-2017

Sample Name	Location	Wheat Variety	Library Name ¹	ENA Run Accession Lane 1 ²	ENA Run Accession Lane 2 ²	Total Read Count	%Mapped to BR32 ³
RI-1	Terial, Kasundi, Gangni	Prodip	LIB27426	ERR1954072 (Reads 1, Reads 2)	ERR1954087 (Reads 1, Reads 2)	40,125,456	0.10%
RI-2	Terial, Kasundi, Gangni	Prodip	LIB27427	ERR1954073 (Reads 1, Reads 2)	ERR1954088 (Reads 1, Reads 2)	43,419,854	0.80%
RI-3	Saharvati, EbadhatKhana, Meherpur	Prodip	LIB27428	ERR1954074 (Reads 1, Reads 2)	ERR1954089 (Reads 1, Reads 2)	39,777,310	0.90%
RI-4	Dhola, Gangni, Meherpur	Prodip	LIB27429	ERR1954075 (Reads 1, Reads 2)	ERR1954090 (Reads 1, Reads 2)	41,585,348	0.40%
RI-5	Garabaria, Gangni, Meherpur	Sotabdi	LIB27430	ERR1954076 (Reads 1, Reads 2)	ERR1954091 (Reads 1, Reads 2)	36,692,872	0.30%
RI-6	Garabaria, Gangni, Meherpur	Sotabdi	LIB27431	ERR1954077 (Reads 1, Reads 2)	ERR1954092 (Reads 1, Reads 2)	42,877,050	0.50%
RI-7	JadupurField, MaherpurSador	BARI gom-26	LIB27432	ERR1954078 (Reads 1, Reads 2)	ERR1954093 (Reads 1, Reads 2)	37,047,832	0.60%
RI-8	JadupurField, MaherpurSador	BARI gom-26	LIB27433	ERR1954079 (Reads 1, Reads 2)	ERR1954094 (Reads 1, Reads 2)	39,404,214	1.60%
RI-9	JadupurField , MaherpurSadar	Prodip	LIB27434	ERR1954080 (Reads 1, Reads 2)	ERR1954095 (Reads 1, Reads 2)	27,380,134	0.90%
RI-10	Dosomi, Kutubpur, Sadar	Sotabdi	LIB27435	ERR1954081 (Reads 1, Reads 2)	ERR1954096 (Reads 1, Reads 2)	41,812,400	0.30%
RI-11	Dosomi, Kutubpur, Sadar	Sotabdi	LIB27436	ERR1954082 (Reads 1, Reads 2)	ERR1954097 (Reads 1, Reads 2)	35,745,922	0.00%
RI-12	Dosomi, Kutubpur, Sadar	Sotabdi	LIB27437	ERR1954083 (Reads 1, Reads 2)	ERR1954098 (Reads 1, Reads 2)	36,219,310	0.00%
RI-13	Kumari, Alamdanga, Chudanga	Prodip	LIB27438	ERR1954084 (Reads 1, Reads 2)	ERR1954099 (Reads 1, Reads 2)	38,964,660	0.10%
RI-14	Kumara, Alamdanga, Chudanga	Prodip	LIB27439	ERR1954085 (Reads 1, Reads 2)	ERR1954100 (Reads 1, Reads 2)	37,971,306	0.10%
RI-15	MostofapurHajipur, MaguraSador	Unknown	LIB27440	ERR1954086 (Reads 1, Reads 2)	ERR1954101 (Reads 1, Reads 2)	38,928,088	0.00%

¹Illumina sequencing libraries prepared by Earlham Institute (Norwich, UK)

²Sequence reads uploaded to European Nucleotide Archive (ENA) database; Average insert size = 268 bp

³Percent of reads mapped to the genomic sequence of wheat blast fungus *Magnaporthe oryzae* pv. *tritici* isolate BR32 using Tophat