												ı	Norma	alized	d(col)	Confu	sion M	latrix d	of Cla	ssifier													_	1 .0
ACC	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.027	0	0	0	0	0	0	0	0	0	0	0		
BLCA -	0	0.83	0	0.048	0	0	0	0.054	0	0	0	0	0	0	0	0.0096	0.01	0	0	0	0	0	0	0.019	0	0.011	0	0	0	0.0092	0	0		
BRCA	0	0.036	0.97	0.065	0	0	0.11	0	0.019	0.077	0.0092 0	0.034	0	0	0	0	0.031	0	0	0	0	0.03	0.031	0.075	0	0	0	0.049	0	0.046	0	0.062		
CESC	0	0	0	0.71	0	0	0	0.027	0.048	0	0	0	0	0	0	0	0.01	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
) ТОНО		0	0	0	0.86	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
COAD	0	0	0	0	0	0.88	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.47	0	0	0.045	0	0	0	0	0	0		
DLBC C	0	0.012	0	0	0	0	0.78	0	0	0	0	0	0	0	0	0	0.01	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		- 0.8
ESCA D		0	0	0	0	0	0	0.68	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.019	0	0.022	0	0	0	0	0	0		
HNSC E		0.024	0	0.065	0	0	0	0.054	0.85	0	0	0	0	0	0	0	0.042	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
KICH H		0	0	0	0	0	0	0	0	0.92	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
KIRC F	0	0	0	0	0	0	0	0	0	0	0.95	0	0	0	0.013	0	0.01	0	0	0	0	0	0	0	0	0.011	0	0	0	0.018	0	0		
KIRP K		0	0	0	0	0	0	0	0	0	0.0092	0.97	0	0	0.013	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
_AML		0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		- 0.6
_	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
LIHC	0	0	0	0	0.14	0	0	0	0	0	0	0	0	0	0.96	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
True LUAD I		0	0	0	0	0	0	0	0	0	0.0092	0	0	0	0.013	0.91	0.052	0	0	0.11	0	0.01	0	0	0	0.022	0	0	0	0.0092	0	0		
Tr. .USC L	0	0.06	0.0045	0.048	0	0.011	0	0.054	0.076	0	0	0	0	0	0	0.029	0.83	0	0	0	0	0	0	0	0.011	0.011	0	0	0	0.0092	0	0		
MESO L			0		0	0	0	0	0	0	0	0	0	0	0	0	0	0.94	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
	0	0 (0.0045	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0.0092	0	0		
AAD -	0	0	0	0.016	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.83	0	0	0	0.019	0	0.011	0	0	0	0	0	0		- 0.4
CPG P	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.97	0	0	0	0	0	0	0	0	0	0	0		
RAD P		0		0	0	0	0	0	0	0	0	0	0	0	0	0	0							0							0			
READ P	0	0.012	0	0	0	0.098	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.5	0	0	0.022	0	0	0	0	0	0		
		0.012	0.009	0	0	0	0.11	0	0	0	0	0	0	0	0	0	0	0.059	0	0	0	0	0	0.87	0	0	0.032	0	0	0	0	0		
		0			0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.98	0	0	0	0	0	0	0		
STAD S	0	0	0	0	0	0.011	0	0.14	0.0095	0	0	0	0	0	0	0.019	0	0	0	0.056	0	0	0	0	0	0.84	0	0	0	0	0	0		- 0.2
TGCT 9		0		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.97	0	0	0	0	0		
		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.0096	0	0	0	0	0	0	0	0	0	0	0	0.94	0	0	0	0		
T MYHT		0			0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0		
		0.012	0.009	0.048	0	0	0	0	0	0	0.018	0	0	0	0	0.019	0	0	0	0	0	0	0	0	0	0	0	0.0097	0	0.9	0.091	0		
n son		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.91	0		
W/0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.011	0	0	0	0	0	0	0.94		
ر	ACC -	BLCA -	BRCA -	CESC -	CHOL -	COAD -	DLBC -	ESCA -	HNSC -	KICH -	KIRC -	KIRP -	LAML -	- 997	LIHC -	LUAD -	- DSC	MESO -	- 70	PAAD -	PCPG -	PRAD -	READ -	SARC -	SKCM -	STAD -	TGCT -	THCA -	- MYHT	UCEC -	ncs -	MVO		- 0.0
					-	J			_								icted	_											•					