		7			
CELE_Y92H12A.2 CELE_F08A8.3					
CELE_Y110A2AR.2					
	CELE_F47H4.10	CELE_F10C5.1			
C	CELE_F CELE_Y66D12A.17	2pΠ2.0 /	CELE_Y50D7A.7	54G11A.6	
		/ / /		ELE_F47G6.2	
CELE_Y1	10A2AM.3				
CELE_Y71G1	2B 15 CELE_C06E2.3	CELE_Y54	1F5B 4	CELE_Y71G12B.10	
OLLL_ITIO	CELE_Y5	/ ~	CELE_ZK5	550.5	
	CELE_K08E3.7	719271181	CELE_F58F9.7	CELE_F08A8.4	
CELE_C52D10.8			\ \\ CELE	_Y69A2AR.5	
			\ \ \		
CELE_C35B1.1	I licitati i mandi della di pre	otoolygia OFLE Bord	\\\\	CELE_Y46	G5A.21
	Ubiquitin mediated pro	oteolysis CELE_R070	CELE_Y108G3AL.1		I E E00110 0
CELE	_F40G9.3		CELE_1 100G3AL.1		LE_F36H9.3
CELE_C52D10.9				CELE_Y25C1A.13	
CELE_Y82E9	OBR 15		Peroxiso	ma CF	ELE_T08B1.6
OLLE 102L	CELE_K04F1.15		//\		
CELE_W02A11,4		CELE_C52D10.7 CELE_C	C52D10.6		
				CELE_F58A6.1	size
CELE_F44G3.6				CELE_C54G	€10.3
CELE	(44.040.40		CELE_Y73F8	A.24 \	Ŏ
CELE_Y	′41C4A.10 / CELE_C54C8.1	1		CELE_Y57A100	C.6
CELE_T05H4.13	CELE_W03H1.2	CELE_Y105C5B.13		CELE_Y54G11A.5 CELI	E_R09E10.3
_	OLLL WOST 11.2	CE	LE_Y54G11A.13	,EEE_134011A.5	
CELE_T08B1.3					TIT TOOD 4.5
- CELE C20110 7	CELE_Y37H2C.:	)			ELE_T02D1.5
CELE_C30H6.7	3222_1371123	_	CELE_#55B11.1	CELE_ZK550.6	
		CELE_Y66D12A.15		CELE_C47A10.5	
CELE_H3	8K22.5	OLLE_TOOD 12/1.10		OELE_04/A10.5	
CELE_T02G5.8	Pyruvate metabolism	0-1-	\\\\	OFFE 7	)   
		CELE	_Y47D3A.23	CELE_Z	\550.4
CELE COEE 4.0	Mucin type Q-	glycan biosynthesis	Basal transcription factor	rs CELE_R02D3.3	
CELE_C05E4.9			Dasai transcription lacto	CELE_Y11	1B2A.16
CE CELE_R11A5.4	LE_T15D6.2				
0222_1(1)/0.1	CELE VZEDONO	CELE_Y17G7B.3		CELE_Y111B	32A.13
	CELE_Y75B8A.9		ELE_Y116F11B.12	_Y49F6B.1	
	CELE_Y48	36A.12		CELE_R119.6	
CELE_W05	G11.6	CEDE (	C38H2.2		
				CELE_C01F1.1	
CELE_LLC1.3 CELE_W09B6.1 CELE_W04A8.7 CELE_F53G2.7					
CELE_T02G5.7 CELE_Y37E11B.4					
CELE 027H5 3					
CELE_Y39G10AL.3					
CELE_Y46H3A.6 CELE_W03F9.5					