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GENERATED BY OPEN BABEL 2.3.1

REMARK 4 XXXX COMPLIES WITH FORMAT V. 2.0

[illegible]

ATOM	932	CB	VAL	100	41.564	56.399	12.933	1.00	0.00	C
ATOM	933	CG1	VAL	100	41.361	56.789	14.408	1.00	0.00	C
ATOM	934	CG2	VAL	100	40.271	56.030	12.186	1.00	0.00	C
ATOM	958	HN	TRP	103	43.676	60.371	12.390	1.00	0.00	H
ATOM	960	CB	TRP	103	42.044	61.476	14.036	1.00	0.00	C
ATOM	961	CG	TRP	103	41.243	62.222	15.086	1.00	0.00	C
ATOM	962	CD2	TRP	103	39.840	62.522	14.961	1.00	0.00	C
ATOM	963	CD1	TRP	103	41.641	62.753	16.280	1.00	0.00	C
ATOM	964	NE1	TRP	103	40.577	63.354	16.908	1.00	0.00	N
ATOM	965	HE1	TRP	103	40.606	63.818	17.816	1.00	0.00	H
ATOM	966	CE2	TRP	103	39.464	63.224	16.106	1.00	0.00	C
ATOM	967	CE3	TRP	103	38.943	62.250	13.966	1.00	0.00	C
ATOM	968	CZ2	TRP	103	38.181	63.665	16.273	1.00	0.00	C
ATOM	969	CZ3	TRP	103	37.647	62.684	14.145	1.00	0.00	C
ATOM	970	CH2	TRP	103	37.273	63.379	15.276	1.00	0.00	C
ATOM	974	HN	CYS	104	44.098	60.276	15.485	1.00	0.00	H
ATOM	977	SG	CYS	104	43.034	59.991	18.433	1.00	0.00	S
ATOM	1484	O	GLY	154	39.615	63.856	11.540	1.00	0.00	O
ATOM	1511	CG1	ILE	157	40.601	62.216	9.603	1.00	0.00	C
ATOM	1513	CD1	ILE	157	40.132	61.087	8.682	1.00	0.00	C
ATOM	7461	CG	GLU	785	32.864	52.846	17.084	1.00	0.00	C
ATOM	7462	CD	GLU	785	32.546	53.684	15.861	1.00	0.00	C
ATOM	7463	OE1	GLU	785	31.649	54.561	15.981	1.00	0.00	O
ATOM	7464	OE2	GLU	785	33.177	53.459	14.794	1.00	0.00	O
ATOM	7502	CG	LEU	789	34.718	55.544	19.754	1.00	0.00	C
ATOM	7504	CD2	LEU	789	34.109	56.288	18.559	1.00	0.00	C
ATOM	8091	CA	PHE	848	40.034	54.565	17.786	1.00	0.00	C
ATOM	8092	CB	PHE	848	40.010	56.108	17.773	1.00	0.00	C
ATOM	8093	CG	PHE	848	39.611	56.579	19.131	1.00	0.00	C
ATOM	8094	CD1	PHE	848	38.285	56.620	19.496	1.00	0.00	C
ATOM	8095	CD2	PHE	848	40.556	56.968	20.049	1.00	0.00	C
ATOM	8096	CE1	PHE	848	37.914	57.048	20.749	1.00	0.00	C
ATOM	8097	CE2	PHE	848	40.195	57.397	21.303	1.00	0.00	C
ATOM	8099	C	PHE	848	40.144	54.093	16.385	1.00	0.00	C
ATOM	8100	O	PHE	848	39.140	53.790	15.742	1.00	0.00	O
ATOM	8119	OG1	THR	851	37.208	52.272	17.094	1.00	0.00	O
ATOM	8120	HG1	THR	851	36.849	52.634	16.292	1.00	0.00	H
ATOM	8124	N	PHE	852	37.143	51.930	14.382	1.00	0.00	N
ATOM	8125	HN	PHE	852	37.851	52.459	14.891	1.00	0.00	H
ATOM	8126	CA	PHE	852	36.470	52.549	13.268	1.00	0.00	C
ATOM	8127	CB	PHE	852	37.019	53.965	12.999	1.00	0.00	C
ATOM	8128	CG	PHE	852	36.596	54.471	11.660	1.00	0.00	C
ATOM	8129	CD1	PHE	852	35.275	54.719	11.366	1.00	0.00	

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