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REM ----- 2PHY
REM 2PHY
REM STRIDE: Knowledge-based secondary structure assignment 2PHY
REM Please cite: D.Frishman & P.Argos, Proteins XX, XXX-XXX, 1995 2PHY
REM 2PHY
REM Residue accessible surface area calculation 2PHY
REM Please cite: F.Eisenhaber & P.Argos, J.Comp.Chem. 14, 1272-1280, 1993 2PHY
REM F.Eisenhaber et al., J.Comp.Chem., 1994, submitted 2PHY
REM 2PHY
REM ----- General information ----- 2PHY
REM 2PHY
HDR PHOTORECEPTOR 12-APR-95 2PHY 2PHY
CMP MOL_ID: 1; 2PHY
CMP MOLECULE: PHOTOACTIVE YELLOW PROTEIN; 2PHY
CMP CHAIN: A; 2PHY
CMP SYNONYM: PYP 2PHY
SRC MOL_ID: 1; 2PHY
SRC ORGANISM_SCIENTIFIC: HALORHODOSPIRA HALOPHILA; 2PHY
SRC ORGANISM_TAXID: 1053; 2PHY
SRC STRAIN: BN9626 2PHY
AUT G.E.O.BORGSTAHL,E.D.GETZOFF 2PHY
REM 2PHY
REM ----- Secondary structure summary ----- 2PHY
REM 2PHY
CHN /home/proj/stride/tmp/tmpR5MTpzpdb A 2PHY
REM 2PHY
REM 2PHY
SEQ 1 MEHVAFGSEDIENTLAKMDDGQLDGLAFGAIQLDGDGNILQYNAAEGDIT 50 2PHY
STR TTTT HHHHHHH HHHHH EEEEEETTTEEEEE HHHHHHH 2PHY
REM 2PHY
REM 2PHY
SEQ 51 GRDPKQVIGKNFFKDVAPCTDSPEFYGKFKEGVASGNLNTMFEYTFDYQM 100 2PHY
STR GGGTTTTEIIIII GGGTTTTTHHHHHHHHHH EEEEEETTTT 2PHY
REM 2PHY
REM 2PHY
SEQ 101 TPTKVKVHMKKALSGDSYWVFVKRV 125 2PHY
STR EEEEEEEE TTTEEEEEEE 2PHY
REM 2PHY
REM 2PHY
REM 2PHY
LOC AlphaHelix ASP 10 A ALA 16 A 2PHY
LOC AlphaHelix ASP 20 A ASP 24 A 2PHY
LOC AlphaHelix ALA 44 A THR 50 A 2PHY
LOC AlphaHelix TYR 76 A SER 85 A 2PHY
LOC 310Helix PRO 54 A GLN 56 A 2PHY
LOC 310Helix PRO 68 A THR 70 A 2PHY
LOC PiHelix PHE 62 A VAL 66 A 2PHY
LOC Strand GLY 29 A ASP 34 A 2PHY
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LOC	Strand	ASN	38	A	TYR	42	A	2PHY
LOC	Strand	LYS	60	A	ASN	61	A	2PHY
LOC	Strand	ASN	89	A	PHE	96	A	2PHY
LOC	Strand	THR	103	A	LYS	111	A	2PHY
LOC	Strand	SER	117	A	ARG	124	A	2PHY
LOC	TurnII	ALA	5	A	SER	8	A	2PHY
LOC	TurnI	ASP	34	A	GLY	37	A	2PHY
LOC	TurnII	VAL	57	A	LYS	60	A	2PHY
LOC	TurnI	SER	72	A	PHE	75	A	2PHY
LOC	TurnIV	PRO	73	A	TYR	76	A	2PHY
LOC	TurnIV	ASP	97	A	MET	100	A	2PHY
LOC	TurnIV	SER	114	A	SER	117	A	2PHY
LOC	GammaInv	THR	70	A	SER	72	A	2PHY
REM								2PHY
REM	----- Detailed secondary structure assignment-----							2PHY
REM								2PHY
REM	---Residue---	--Structure--		-Phi-		-Psi-		2PHY
ASG	MET A	1	1	C	Coil	360.00	146.81	190.6
ASG	GLU A	2	2	C	Coil	-82.98	126.40	45.0
ASG	HIS A	3	3	C	Coil	-90.49	85.97	169.8
ASG	VAL A	4	4	C	Coil	-138.24	132.92	16.7
ASG	ALA A	5	5	T	Turn	-91.24	137.96	60.6
ASG	PHE A	6	6	T	Turn	-55.73	131.36	33.8
ASG	GLY A	7	7	T	Turn	84.75	5.75	30.6
ASG	SER A	8	8	T	Turn	-59.42	138.95	40.7
ASG	GLU A	9	9	C	Coil	-59.57	-31.59	162.3
ASG	ASP A	10	10	H	AlphaHelix	-109.87	26.30	94.3
ASG	ILE A	11	11	H	AlphaHelix	-54.94	-38.42	0.4
ASG	GLU A	12	12	H	AlphaHelix	-62.43	-33.37	37.6
ASG	ASN A	13	13	H	AlphaHelix	-71.42	-42.81	91.8
ASG	THR A	14	14	H	AlphaHelix	-63.86	-42.99	42.0
ASG	LEU A	15	15	H	AlphaHelix	-77.88	-14.85	2.3
ASG	ALA A	16	16	H	AlphaHelix	-61.52	-24.49	68.1
ASG	LYS A	17	17	C	Coil	-88.83	-5.23	136.6
ASG	MET A	18	18	C	Coil	-93.80	124.52	12.1
ASG	ASP A	19	19	C	Coil	-76.47	170.85	84.2
ASG	ASP A	20	20	H	AlphaHelix	-57.24	-32.22	76.9
ASG	GLY A	21	21	H	AlphaHelix	-67.33	-37.22	39.4
ASG	GLN A	22	22	H	AlphaHelix	-75.82	-33.38	67.3
ASG	LEU A	23	23	H	AlphaHelix	-61.57	-32.44	1.6
ASG	ASP A	24	24	H	AlphaHelix	-67.68	-15.73	67.4
ASG	GLY A	25	25	C	Coil	-89.78	2.50	27.3
ASG	LEU A	26	26	C	Coil	-72.79	151.36	8.4
ASG	ALA A	27	27	C	Coil	-79.58	-1.84	65.8
ASG	PHE A	28	28	C	Coil	-141.34	165.84	6.7
ASG	GLY A	29	29	E	Strand	-71.86	141.94	0.0
ASG	ALA A	30	30	E	Strand	-143.22	127.89	0.0
ASG	ILE A	31	31	E	Strand	-128.84	132.59	0.2
ASG	GLN A	32	32	E	Strand	-110.50	128.64	26.9

ASG	LEU	A	33	33	E	Strand	-118.83	152.07	0.0	2PHY
ASG	ASP	A	34	34	E	Strand	-89.38	-178.19	38.5	2PHY
ASG	GLY	A	35	35	T	Turn	-60.60	-21.22	19.9	2PHY
ASG	ASP	A	36	36	T	Turn	-92.97	-1.48	120.7	2PHY
ASG	GLY	A	37	37	T	Turn	92.47	10.11	0.0	2PHY
ASG	ASN	A	38	38	E	Strand	-79.87	139.92	48.7	2PHY
ASG	ILE	A	39	39	E	Strand	-81.21	121.13	1.7	2PHY
ASG	LEU	A	40	40	E	Strand	-100.36	-35.39	78.8	2PHY
ASG	GLN	A	41	41	E	Strand	-142.44	149.95	33.1	2PHY
ASG	TYR	A	42	42	E	Strand	-157.65	112.67	12.5	2PHY
ASG	ASN	A	43	43	C	Coil	-85.53	169.75	0.4	2PHY
ASG	ALA	A	44	44	H	AlphaHelix	-61.61	-40.64	35.0	2PHY
ASG	ALA	A	45	45	H	AlphaHelix	-61.22	-35.09	17.2	2PHY
ASG	GLU	A	46	46	H	AlphaHelix	-67.82	-41.96	2.0	2PHY
ASG	GLY	A	47	47	H	AlphaHelix	-63.44	-33.23	8.6	2PHY
ASG	ASP	A	48	48	H	AlphaHelix	-66.96	-38.26	142.0	2PHY
ASG	ILE	A	49	49	H	AlphaHelix	-73.47	-39.47	41.9	2PHY
ASG	THR	A	50	50	H	AlphaHelix	-102.85	-17.57	13.2	2PHY
ASG	GLY	A	51	51	C	Coil	91.01	-12.64	67.6	2PHY
ASG	ARG	A	52	52	C	Coil	-71.90	147.71	45.3	2PHY
ASG	ASP	A	53	53	C	Coil	-89.56	119.88	84.6	2PHY
ASG	PRO	A	54	54	G	310Helix	-47.42	-52.51	31.7	2PHY
ASG	LYS	A	55	55	G	310Helix	-61.81	-27.56	133.8	2PHY
ASG	GLN	A	56	56	G	310Helix	-79.86	-23.23	133.9	2PHY
ASG	VAL	A	57	57	T	Turn	-94.06	-14.84	9.8	2PHY
ASG	ILE	A	58	58	T	Turn	-63.96	129.23	66.4	2PHY
ASG	GLY	A	59	59	T	Turn	99.42	-16.02	58.3	2PHY
ASG	LYS	A	60	60	E	Strand	-83.39	158.43	92.9	2PHY
ASG	ASN	A	61	61	E	Strand	-86.42	123.37	25.4	2PHY
ASG	PHE	A	62	62	I	PiHelix	-60.08	-46.31	7.0	2PHY
ASG	PHE	A	63	63	I	PiHelix	-78.87	-30.37	2.7	2PHY
ASG	LYS	A	64	64	I	PiHelix	-92.89	-43.82	163.7	2PHY
ASG	ASP	A	65	65	I	PiHelix	-95.54	-57.28	80.7	2PHY
ASG	VAL	A	66	66	I	PiHelix	-79.91	-42.77	16.3	2PHY
ASG	ALA	A	67	67	C	Coil	-129.57	66.76	11.5	2PHY
ASG	PRO	A	68	68	G	310Helix	-59.50	-25.75	47.0	2PHY
ASG	CYS	A	69	69	G	310Helix	-66.34	-17.47	31.2	2PHY
ASG	THR	A	70	70	G	310Helix	-91.32	-6.20	0.2	2PHY
ASG	ASP	A	71	71	T	Turn	-80.70	85.03	93.9	2PHY
ASG	SER	A	72	72	T	Turn	-161.02	165.37	31.6	2PHY
ASG	PRO	A	73	73	T	Turn	-62.54	-27.21	111.6	2PHY
ASG	GLU	A	74	74	T	Turn	-75.49	-13.94	125.7	2PHY
ASG	PHE	A	75	75	T	Turn	-130.59	-74.27	0.0	2PHY
ASG	TYR	A	76	76	H	AlphaHelix	-55.56	-39.15	83.8	2PHY
ASG	GLY	A	77	77	H	AlphaHelix	-59.77	-38.15	12.5	2PHY
ASG	LYS	A	78	78	H	AlphaHelix	-71.78	-38.16	60.8	2PHY
ASG	PHE	A	79	79	H	AlphaHelix	-63.34	-49.53	2.4	2PHY
ASG	LYS	A	80	80	H	AlphaHelix	-66.27	-30.48	125.4	2PHY
ASG	GLU	A	81	81	H	AlphaHelix	-68.32	-41.07	128.8	2PHY

ASG	GLY	A	82	82	H	AlphaHelix	-67.59	-41.24	4.3	2PHY
ASG	VAL	A	83	83	H	AlphaHelix	-63.19	-43.51	32.3	2PHY
ASG	ALA	A	84	84	H	AlphaHelix	-66.34	-36.01	90.9	2PHY
ASG	SER	A	85	85	H	AlphaHelix	-79.83	-21.94	96.0	2PHY
ASG	GLY	A	86	86	C	Coil	81.59	14.79	37.4	2PHY
ASG	ASN	A	87	87	C	Coil	-151.34	138.01	118.0	2PHY
ASG	LEU	A	88	88	C	Coil	-159.21	128.29	12.7	2PHY
ASG	ASN	A	89	89	E	Strand	-163.00	94.56	97.6	2PHY
ASG	THR	A	90	90	E	Strand	-145.12	147.14	42.9	2PHY
ASG	MET	A	91	91	E	Strand	-131.41	134.71	76.4	2PHY
ASG	PHE	A	92	92	E	Strand	-159.91	159.51	28.1	2PHY
ASG	GLU	A	93	93	E	Strand	-89.54	150.12	123.7	2PHY
ASG	TYR	A	94	94	E	Strand	-152.42	160.51	34.3	2PHY
ASG	THR	A	95	95	E	Strand	-115.61	121.51	56.9	2PHY
ASG	PHE	A	96	96	E	Strand	-91.03	123.74	11.6	2PHY
ASG	ASP	A	97	97	T	Turn	-141.79	18.54	86.7	2PHY
ASG	TYR	A	98	98	T	Turn	-92.02	116.57	94.7	2PHY
ASG	GLN	A	99	99	T	Turn	56.98	33.56	156.0	2PHY
ASG	MET	A	100	100	T	Turn	-159.39	165.73	42.7	2PHY
ASG	THR	A	101	101	C	Coil	-67.90	131.38	115.3	2PHY
ASG	PRO	A	102	102	C	Coil	-52.09	122.51	91.6	2PHY
ASG	THR	A	103	103	E	Strand	-124.57	121.80	16.5	2PHY
ASG	LYS	A	104	104	E	Strand	-80.16	129.34	131.2	2PHY
ASG	VAL	A	105	105	E	Strand	-136.69	162.96	0.0	2PHY
ASG	LYS	A	106	106	E	Strand	-106.67	128.82	68.1	2PHY
ASG	VAL	A	107	107	E	Strand	-114.79	135.25	0.0	2PHY
ASG	HIS	A	108	108	E	Strand	-124.85	116.58	4.0	2PHY
ASG	MET	A	109	109	E	Strand	-106.04	129.90	0.0	2PHY
ASG	LYS	A	110	110	E	Strand	-147.61	138.72	21.2	2PHY
ASG	LYS	A	111	111	E	Strand	-64.70	135.54	57.8	2PHY
ASG	ALA	A	112	112	C	Coil	-74.85	167.78	5.6	2PHY
ASG	LEU	A	113	113	C	Coil	-67.87	-34.52	100.7	2PHY
ASG	SER	A	114	114	T	Turn	-63.01	162.77	76.8	2PHY
ASG	GLY	A	115	115	T	Turn	-57.32	-74.29	51.8	2PHY
ASG	ASP	A	116	116	T	Turn	-107.86	58.81	114.4	2PHY
ASG	SER	A	117	117	E	Strand	-141.28	147.95	4.2	2PHY
ASG	TYR	A	118	118	E	Strand	-137.62	144.12	2.4	2PHY
ASG	TRP	A	119	119	E	Strand	-101.33	146.83	5.9	2PHY
ASG	VAL	A	120	120	E	Strand	-127.19	115.18	0.2	2PHY
ASG	PHE	A	121	121	E	Strand	-107.74	143.29	0.2	2PHY
ASG	VAL	A	122	122	E	Strand	-140.35	137.11	0.2	2PHY
ASG	LYS	A	123	123	E	Strand	-143.37	153.53	84.2	2PHY
ASG	ARG	A	124	124	E	Strand	-71.17	148.60	183.9	2PHY
ASG	VAL	A	125	125	C	Coil	-130.63	360.00	102.2	2PHY