_	Pactoroidatas Ignovihantaria: Kryntonialas: PSV/26	Chloroflevi: Debalesessoidis: H2 02	Dianharatritas: Miararahasia	Furvershapeta: Thermonlasmeta: uncultured: uncultured erebasen	Commetimenedates: MD2002, B12: ungultured besterium: ungultured besterium	Nanograhagagata: Wagagarahagia: Candidatus Daggarahagata arahagan BBC 10ET COMBO 24 0: Candidatus Daggarahagata arahagan BBC 10ET COMBO 24 0	Potosoibactorio: APV1: Candidatus Falkowbactorio: unaulturad bactorium	Petagoihagtaria: CRR2: ungulturad hagtarium: ungulturad hagtarium	Potoscibactorio: Migroganometio: Candidatus Wassahactoria: Ambiguous, taya: Ambiguous, taya: Ambiguous, taya	Pataggibagteria: Pargubagteria: Candidatus Stackowiezbagteria	Planetemyostos: Physianhaerae: S. 70	Protechectoria: Alphanrotachactoria: Sphingemanadales: Sphingemanadaeses	Protophostoria: Poltoprotophostoria: Muyaccondos: P2OP, 42	Proteobacteria; Gammaproteobacteria; Cellvibrionales; Porticoccace
														Proteobacteria; Gammaproteobacteria; Coxiellales; Coxiellaceae
														Proteobacteria; Gammaproteobacteria; Ectothiorhodospirales; Ectothi
	BRC1	Chloroflexi; Dehalococcoidia; SAR202 clade	Edwardsbacteria; uncultured bacterium; uncultured bacterium; uncultured bacterium	Fibrobacteres; Fibrobacteria; Fibrobacterales; Candidatus Raymondbacteria	Kiritimatiellaeota; Kiritimatiellae; WCHB1-41	Nanoarchaeaeota; Woesearchaeia; uncultured euryarchaeote; uncultured euryarchaeote	Patescibacteria; ABY1; Candidatus Komeilibacteria	Patescibacteria; Gracilibacteria; Candidatus Peregrinibacteria; uncultured bacterium	Patescibacteria; Microgenomatia; Candidatus Woesebacteria; metagenome	Patescibacteria; Parcubacteria; Candidatus Yanofskybacteria; uncultured bacterium	Planctomycetes; Planctomycetacia; Pirellulales; Pirellulaceae	Proteobacteria; Deltaproteobacteria; Bdellovibrionales; Bdellovibrionaceae	Proteobacteria; Deltaproteobacteria; PB19; uncultured organism	Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobact
	Caldiserica; Caldisericales; TTA-B15	Chloroflexi; Dehalococcoidia; SAR202 clade; uncultured bacterium	Elusimicrobia; Elusimicrobia; Lineage IV	Firestonebacteria	Latescibacteria	Nitrospinae; P9X2b3D02	Patescibacteria; ABY1; Candidatus Komeilibacteria; Parcubacteria group bacterium GW2011_GWF2_45_11	Patescibacteria; Gracilibacteria; Candidatus Peregrinibacteria; uncultured marine bacterium	Patescibacteria; Microgenomatia; Candidatus Woykebacteria; Candidatus Woykebacteria bacterium RBG_13_40_7b	Patescibacteria; Parcubacteria; Candidatus Zambryskibacteria	Planctomycetes; SPG12–343–353–B69; uncultured bacterium; uncultured bacteriun	Proteobacteria; Deltaproteobacteria; Deltaproteobacteria Incertae Sedis; Unknown Famil	Proteobacteria; Deltaproteobacteria; SAR324 clade(Marine group B)	Proteobacteria; Gammaproteobacteria; Gammaproteobacteria Incerta
	Calditrichaeota; Calditrichia; Calditrichales; Calditrichaceae	Chloroflexi; Dehalococcoidia; SAR202 clade; uncultured eubacterium H1.2.f	Elusimicrobia; Elusimicrobia; Lineage IV; uncultured bacterium	Firmicutes	Latescibacteria; Latescibacterales; Latescibacteraceae	Nitrospirae; 4–29–1	Patescibacteria; ABY1; Candidatus Kuenenbacteria	Patescibacteria; Gracilibacteria; Candidatus Peribacteria	Patescibacteria; Parcubacteria	Patescibacteria; Parcubacteria; CG1-02-42-13; Parcubacteria group bacterium CG1_02_42_	3 Planctomycetes; vadinHA49	Proteobacteria; Deltaproteobacteria; Desulfarculales; Desulfarculaceae	Proteobacteria; Deltaproteobacteria; Sva0485	Proteobacteria; Gammaproteobacteria; Halothiobacillales; Halothiobac
acteroidales	Chlamydiae; Chlamydiales	Chloroflexi; Dehalococcoidia; Sh765B-AG-111; uncultured bacterium	Elusimicrobia; Lineage IIc	Firmicutes; Bacilli; Lactobacillales; Enterococcaceae	Latescibacteria; uncultured bacterium; uncultured bacterium; uncultured bacterium	Nitrospirae; 4–29–1; uncultured bacterium; uncultured bacterium	Patescibacteria; ABY1; Candidatus Magasanikbacteria	Patescibacteria; Microgenomatia; Candidatus Amesbacteria; Candidatus Amesbacteria bacterium GW2011_GWA2_42_12	Patescibacteria; Parcubacteria; Candidatus Azambacteria	Patescibacteria; Parcubacteria; Paceibacterales; Paceibacteraceae	Planctomycetes; vadinHA49; uncultured bacterium; uncultured bacterium	Proteobacteria; Deltaproteobacteria; Desulfobacterales; Desulfobacteraceae	Proteobacteria; Deltaproteobacteria; Sva0485; uncultured delta proteobacterium	Proteobacteria; Gammaproteobacteria; Legionellales; Legionellaceae
acteroidales; Bacteroidaceae	Chlamydiae; Chlamydiales; cvE6	Chloroflexi; TK10	Elusimicrobia; Lineage IIc; Elusimicrobia bacterium RIFOXYA2_FULL_39_19; Elusimicrobia bacterium RIFOXYA2_FULL_39_19	Firmicutes; Clostridia	Latescibacteria; uncultured organism; uncultured organism; uncultured organism	Nitrospirae; Nitrospira; Nitrospirales; Nitrospiraceae	Patescibacteria; ABY1; Candidatus Magasanikbacteria; Candidatus Magasanikbacteria bacterium GW2011_GWA2_41_55	Patescibacteria; Microgenomatia; Candidatus Amesbacteria; Microgenomates bacterium SCGC AAA011-B20	Patescibacteria; Parcubacteria; Candidatus Azambacteria; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa	Patescibacteria; Parcubacteria; UBA9983	Planctomycetes; vadinHA49; uncultured organism; uncultured organism	Proteobacteria; Deltaproteobacteria; Desulfobacterales; Desulfobulbaceae	Proteobacteria; Deltaproteobacteria; Syntrophobacterales	Proteobacteria; Gammaproteobacteria; Methylococcales; Methylomona
acteroidales; Bacteroidetes BD2-2	Chlamydiae; Chlamydiales; Parachlamydiaceae	Crenarchaeota; Bathyarchaeia	Elusimicrobia; Rs-M47; uncultured bacterium; uncultured bacterium	Firmicutes; Clostridia; Clostridiales; Christensenellaceae	LCP-89; uncultured bacterium; uncultured bacterium; uncultured bacterium	Nitrospirae; Thermodesulfovibrionia; uncultured	Patescibacteria; ABY1; Candidatus Magasanikbacteria; Candidatus Magasanikbacteria bacterium GW2011_GWC2_45_8	Patescibacteria; Microgenomatia; Candidatus Beckwithbacteria; uncultured bacterium	Patescibacteria; Parcubacteria; Candidatus Azambacteria; uncultured bacterium	Patescibacteria; Parcubacteria; uncultured bacterium; uncultured bacterium	Poribacteria	Proteobacteria; Deltaproteobacteria; Desulfovibrionales; Desulfomicrobiaceae	Proteobacteria; Deltaproteobacteria; Syntrophobacterales; Syntrophaceae	Proteobacteria; Gammaproteobacteria; Pseudomonadales; Moraxellac
acteroidales; Bacteroidetes vadinHA17	Chlamydiae; LD1-PA32	Crenarchaeota; Bathyarchaeia; uncultured archaeon; uncultured archaeon	Elusimicrobia; Rs-M47; uncultured organism; uncultured organism	Firmicutes; Clostridia; Clostridiales; Eubacteriaceae	Lentisphaerae	Nitrospirae; Thermodesulfovibrionia; uncultured; uncultured bacterium	Patescibacteria; ABY1; Candidatus Magasanikbacteria; Candidatus Magasanikbacteria bacterium RIFCSPHIGHO2_12_FULL_41_16	Patescibacteria; Microgenomatia; Candidatus Collierbacteria	Patescibacteria; Parcubacteria; Candidatus Colwellbacteria	Patescibacteria; Parcubacteria; uncultured organism; uncultured organism	Proteobacteria; Alphaproteobacteria	Proteobacteria; Deltaproteobacteria; Desulfovibrionales; Desulfovibrionaceae	Proteobacteria; Deltaproteobacteria; Syntrophobacterales; Syntrophobacteraceae	Proteobacteria; Gammaproteobacteria; Salinisphaerales; Solimonada
acteroidales; Prolixibacteraceae	Chlamydiae; LD1-PA32; uncultured bacterium; uncultured bacterium	Cyanobacteria; Melainabacteria; Caenarcaniphilales; uncultured bacterium	Epsilonbacteraeota; Campylobacteria; Campylobacterales; Arcobacteraceae	Firmicutes; Clostridia; Clostridiales; Family XVIII	Lentisphaerae; Lentisphaeria; Victivallales	Omnitrophicaeota	Patescibacteria; ABY1; Candidatus Magasanikbacteria; Candidatus Magasanikbacteria bacterium RIFOXYD2_FULL_36_9	Patescibacteria; Microgenomatia; Candidatus Collierbacteria; groundwater metagenome	Patescibacteria; Parcubacteria; Candidatus Colwellbacteria; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa	Patescibacteria; WWE3	Proteobacteria; Alphaproteobacteria; Caulobacterales; Caulobacteraceae	Proteobacteria; Deltaproteobacteria; Desulfuromonadales	Proteobacteria; Gammaproteobacteria	Proteobacteria; Gammaproteobacteria; Thiotrichales; Thiotrichaceae
acteroidales; Rikenellaceae	Chloroflexi; Anaerolineae	Cyanobacteria; Melainabacteria; Gastranaerophilales	Epsilonbacteraeota; Campylobacteria; Campylobacterales; Sulfurovaceae	Firmicutes; Clostridia; Clostridiales; Gracilibacteraceae	Lentisphaerae; Lentisphaeria; Victivallales; GWF2-44-16	Omnitrophicaeota; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa	Patescibacteria; ABY1; Candidatus Magasanikbacteria; Candidatus Uhrbacteria bacterium RIFCSPHIGHO2_12_FULL_60_25	Patescibacteria; Microgenomatia; Candidatus Curtissbacteria; uncultured bacterium	Patescibacteria; Parcubacteria; Candidatus Jorgensenbacteria; Candidatus Adlerbacteria bacterium GW2011_GWC1_50_9	PAUC34f	Proteobacteria; Alphaproteobacteria; Caulobacterales; Hyphomonadaceae	Proteobacteria; Deltaproteobacteria; DTB120	Proteobacteria; Gammaproteobacteria; Acidiferrobacterales; Acidiferrobacteraceae	Proteobacteria; Gammaproteobacteria; Xanthomonadales; Xanthomo
hitinophagales; Chitinophagaceae	Chloroflexi; Anaerolineae; ADurb.Bin180	Cyanobacteria; Melainabacteria; Obscuribacterales	Euryarchaeota; Methanobacteria; Methanobacteriales; Methanobacteriaceae	Firmicutes; Clostridia; Clostridiales; Peptococcaceae	Lentisphaerae; Lentisphaeria; Victivallales; PRD18C08	Omnitrophicaeota; Omnitrophia; Omnitrophales; Omnitrophaceae	Patescibacteria; ABY1; Candidatus Magasanikbacteria; uncultured bacterium	Patescibacteria; Microgenomatia; Candidatus Daviesbacteria	Patescibacteria; Parcubacteria; Candidatus Kaiserbacteria	Planctomycetes; Brocadiae; Brocadiales; Brocadiaceae	Proteobacteria; Alphaproteobacteria; Paracaedibacterales; Paracaedibacteraceae	Proteobacteria; Deltaproteobacteria; DTB120; uncultured bacterium	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales	Rokubacteria; NC10; Methylomirabilales; Methylomirabilaceae
lavobacteriales; Crocinitomicaceae	Chloroflexi; Anaerolineae; Anaerolineales; Anaerolineaceae	Dependentiae; Babeliae; Babeliales	Euryarchaeota; Methanomicrobia; Methanosarcinales; Methanoperedenaceae	Firmicutes; Clostridia; Clostridiales; Ruminococcaceae	Lentisphaerae; Oligosphaeria; Oligosphaerales; Lenti-02	Omnitrophicaeota; Omnitrophica WOR_2 bacterium RIFCSPLOWO2_12_FULL_51_24; Omnitrophica WOR_2 bacterium RIFCSPLOWO2_12_FULL_51_24; Omnitrophica WOR_2 bacterium RIFCSPLOWO2_12_FULL_51_24	Patescibacteria; ABY1; Candidatus Magasanikbacteria; uncultured Parcubacteria group bacterium	Patescibacteria; Microgenomatia; Candidatus Daviesbacteria; uncultured bacterium	Patescibacteria; Parcubacteria; Candidatus Kaiserbacteria; metagenome	Planctomycetes; OM190	Proteobacteria; Alphaproteobacteria; Parvibaculales; Parvibaculaceae	Proteobacteria; Deltaproteobacteria; MBNT15	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Burkholderiaceae	Rokubacteria; NC10; Rokubacteriales; uncultured bacterium
phingobacteriales	Chloroflexi; Anaerolineae; RBG-13-54-9	Desantisbacteria; Candidatus Desantisbacteria bacterium CG1_02_49_89; Candidatus Desantisbacteria bacterium CG1_02_49_89; Candidatus Desantisbacteria bacterium CG1_02_49_89	Euryarchaeota; Methanomicrobia; Methanosarcinales; Methanosarcinaceae	Firmicutes; Clostridia; Clostridiales; Syntrophomonadaceae	Nanoarchaeaeota; Nanohaloarchaeia; Aenigmarchaeales; uncultured archaeon	Omnitrophicaeota; uncultured bacterium; uncultured bacterium uncultured bacterium	Patescibacteria; ABY1; Candidatus Uhrbacteria	Patescibacteria; Microgenomatia; Candidatus Daviesbacteria; uncultured Microgenomates group bacterium	Patescibacteria; Parcubacteria; Candidatus Liptonbacteria	Planctomycetes; Phycisphaerae; CCM11a	Proteobacteria; Alphaproteobacteria; Reyranellales; Reyranellaceae	Proteobacteria; Deltaproteobacteria; Myxococcales	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Gallionellaceae	Schekmanbacteria; uncultured bacterium; uncultured bacterium; uncul
phingobacteriales; Lentimicrobiaceae	Chloroflexi; Anaerolineae; RBG-13-54-9; uncultured bacterium	Diapherotrites; lainarchaeia; lainarchaeales	Euryarchaeota; Thermoplasmata	Firmicutes; Clostridia; D8A-2	Nanoarchaeaeota; Nanohaloarchaeia; Deep Sea Euryarchaeotic Group(DSEG); uncultured archaeon	Omnitrophicaeota; uncultured Verrucomicrobia bacterium; uncultured Verrucomicrobia bacterium; uncultured Verrucomicrobia bacterium	Patescibacteria; ABY1; Candidatus Uhrbacteria; Candidatus Uhrbacteria bacterium GW2011_GWE2_45_35	Patescibacteria; Microgenomatia; Candidatus Gottesmanbacteria	Patescibacteria; Parcubacteria; Candidatus Lloydbacteria	Planctomycetes; Phycisphaerae; DG-20; uncultured bacterium	Proteobacteria; Alphaproteobacteria; Rhizobiales	Proteobacteria; Deltaproteobacteria; Myxococcales; Archangiaceae	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Hydrogenophilacea	Spirochaetes; Leptospirae; Leptospirales; Leptospiraceae
; Ignavibacteriales	Chloroflexi; Anaerolineae; SBR1031	Diapherotrites; lainarchaeia; lainarchaeales; Diapherotrites archaeon SCGC AAA011-K09	Euryarchaeota; Thermoplasmata; Marine Group II	Firmicutes; Clostridia; D8A-2; uncultured bacterium	Nanoarchaeaeota; Woesearchaeia	Patescibacteria; ABY1; Candidatus Buchananbacteria	Patescibacteria; ABY1; Candidatus Uhrbacteria; uncultured bacterium	Patescibacteria; Microgenomatia; Candidatus Levybacteria	Patescibacteria; Parcubacteria; Candidatus Moranbacteria	Planctomycetes; Phycisphaerae; mle1-8	Proteobacteria; Alphaproteobacteria; Rhizobiales; Beijerinckiaceae	Proteobacteria; Deltaproteobacteria; Myxococcales; bacteriap25	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Methylophilaceae	Spirochaetes; MVP–15; uncultured bacterium; uncultured bacterium
		Diapherotrites; lainarchaeia; lainarchaeales; Diapherotrites archaeon SCGC AAA011-N19	Euryarchaeota; Thermoplasmata; Marine Group II; uncultured euryarchaeote	Firmicutes; Clostridia; Thermoanaerobacterales; Thermoanaerobacteraceae	Nanoarchaeaeota; Woesearchaeia; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa; Ambiguous_taxa	Patescibacteria; ABY1; Candidatus Buchananbacteria; Parcubacteria group bacterium GW2011_GWA2_38_13	Patescibacteria; Berkelbacteria	Patescibacteria; Microgenomatia; Candidatus Roizmanbacteria; uncultured bacterium	Patescibacteria; Parcubacteria; Candidatus Moranbacteria; uncultured soil bacterium		Proteobacteria; Alphaproteobacteria; Rhizobiales; Hyphomicrobiaceae	Proteobacteria; Deltaproteobacteria; Myxococcales; Blfdi19	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Nitrosomonadacea	Spirochaetes; Spirochaetia; Spirochaetales; Spirochaetaceae
; Ignavibacteriales; Ignavibacteriaceae	Chloroflexi; Dehalococcoidia; GIF9	Diapherotrites; lainarchaeia; lainarchaeales; uncultured archaeon	Euryarchaeota; Thermoplasmata; Methanomassiliicoccales; uncultured	Firmicutes; Negativicutes; Selenomonadales; Veillonellaceae	Nanoarchaeaeota; Woesearchaeia; archaeon GW2011_AR18; archaeon GW2011_AR18	Patescibacteria; ABY1; Candidatus Buchananbacteria; uncultured bacterium	Patescibacteria; Berkelbacteria; candidate division WS2 bacterium ADurb.Bin280; candidate division WS2 bacterium ADurb.Bin280	Patescibacteria; Microgenomatia; Candidatus Shapirobacteria	Patescibacteria; Parcubacteria; Candidatus Nomurabacteria	Planctomycetes; Phycisphaerae; Phycisphaerales; AKAU3564 sediment group	Proteobacteria; Alphaproteobacteria; Rhizobiales; Rhizobiaceae	Proteobacteria; Deltaproteobacteria; Myxococcales; Haliangiaceae	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Rhodocyclaceae	Thaumarchaeota; Marine Benthic Group A
; Ignavibacteriales; PHOS-HE36	Chloroflexi; Dehalococcoidia; GIF9; uncultured bacterium	Diapherotrites; lainarchaeia; uncultured archaeon; uncultured archaeon	Euryarchaeota; Thermoplasmata; uncultured	Gemmatimonadetes; Gemmatimonadetes; Gemmatimonadales; Gemmatimonadaceae	Nanoarchaeaeota; Woesearchaeia; Candidatus Diapherotrites archaeon ADurb.Bin253; Candidatus Diapherotrites archaeon ADurb.Bin25	Patescibacteria; ABY1; Candidatus Falkowbacteria	Patescibacteria; Berkelbacteria; uncultured bacterium; uncultured bacterium	Patescibacteria; Microgenomatia; Candidatus Woesebacteria	Patescibacteria; Parcubacteria; Candidatus Nomurabacteria; uncultured bacterium	Planctomycetes; Phycisphaerae; Phycisphaerales; Phycisphaeraceae	Proteobacteria; Alphaproteobacteria; Rickettsiales; Mitochondria	Proteobacteria; Deltaproteobacteria; Myxococcales; mle1–27	Proteobacteria; Gammaproteobacteria; Betaproteobacteriales; Sulfuricellaceae	Thaumarchaeota; Nitrososphaeria; Nitrosopumilales; Nitrosopumilac
	ncultured bacterium; uncultured bacterium d  Bacteroidales Bacteroidales; Bacteroidaceae Bacteroidales; Bacteroidetes BD2–2 Bacteroidales; Bacteroidetes vadinHA17 Bacteroidales; Prolixibacteraceae Bacteroidales; Rikenellaceae Chitinophagales; Chitinophagaceae Chitinophagales; Crocinitomicaceae Chitinophagales; Crocinitomicaceae Chingobacteriales	Bacteroidetes; Ignavibacteria; OPB56  ncultured bacterium; uncultured bacterium  Bacteroidetes; Ignavibacteria; SJA–28  BRC1  Caldiserica; Caldisericia; Caldisericales; TTA–B15  Calditrichaeota; Calditrichia; Calditrichaes; Calditrichaceae  Chlamydiae; Chlamydiae; Chlamydiales  Chlamydiae; Chlamydiae; Chlamydiales; cvE6  Cacteroidales; Bacteroidetes BD2–2  Chlamydiae; Chlamydiae; Chlamydiaes; Parachlamydiaceae  Chlamydiae; Chlamydiae; Chlamydiaes; Parachlamydiaceae  Chlamydiae; LD1–PA32  Chlamydiae; LD1–PA32  Chlamydiae; LD1–PA32; uncultured bacterium; uncultured bacterium  Cacteroidales; Rikenellaceae  Chloroflexi; Anaerolineae  Chloroflexi; Anaerolineae; ADurb.Bin180  Chloroflexi; Anaerolineae; Anaerolineales; Anaerolineaceae  Chloroflexi; Anaerolineae; RBG–13–54–9  Chloroflexi; Anaerolineae; RBG–13–54–9  Chloroflexi; Anaerolineae; RBG–13–54–9; uncultured bacterium  Chloroflexi; Anaerolineae; SBR1031  Chloroflexi; Dehalococcoidia  Chloroflexi; Dehalococcoidia; GIF9	Bacteroidetes; Ignavibacteria; OPB56 Chloroflexi; Dehalococcocidia; MSBL5, uncultured bacterium  BRC1 Chloroflexi; Dehalococcodidia; S085, uncultured bacterium  BRC1 Chloroflexi; Dehalococcodidia; S085, uncultured bacterium  BRC1 Chloroflexi; Dehalococcodidia; S085, uncultured bacterium  Caldisrichaeria; Caldisericia, Caldisericiales; TTA-B15 Chloroflexi; Dehalococcodidia; SAR202 clade; uncultured bacterium  Caldirichaeota; Caldisrichaes, Calditrichaeota; Calditrichaeota; Calditrichaeota; Calditrichaeota; Calditrichaeota; Chlamydiales  Chloroflexi; Dehalococcodidia; SAR202 clade; uncultured bacterium  Chloroflexi; Dehalococcodidia; SAR202 clade; uncultured eubacterium H1.2.f.  Caldirichaeota; Chlamydiae; Chlamydiales; Chlamydiales; Chlamydiales; Chlamydiales; Chlamydiales; Chlamydiales; Chlamydiales; Polamydiales; Po	Sectionidates, Irganizationidates, Irganizatio	Descriptions (productions (prod	CPUER CONTINUE NUMBER CONTIN	The Parlies of Table (1995) and Table (1	See And Jackson A 1966.  A 1966 A 196	Medical medica	Printing of the state of the st	Segment of the segmen		See Learning Comment of the Comment	A TOUR AND T