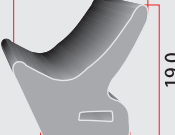
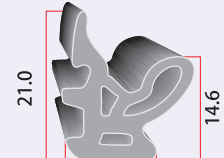
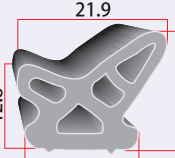
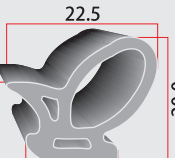
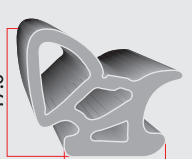
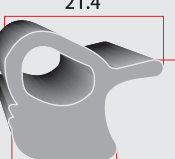
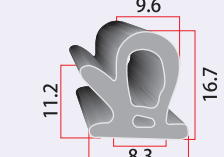
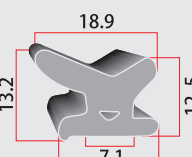
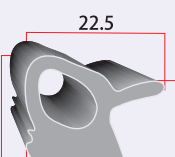
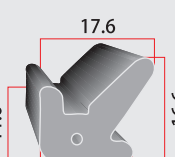
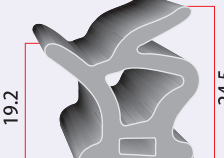
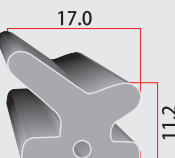
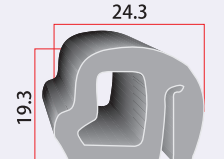
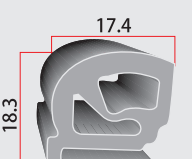
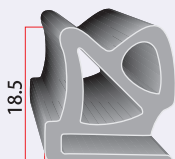




CELLULAR PROFILES | SPONGES

 32.9 20.0 ALP-05005	 18.1 16.4 17.7 16.3 ALP-05010	 18.5 17.5 14.0 ALP-05016	 19.0 19.0 12.0 ALP-07005
 21.0 14.6 12.4 ALP-07106A	 23.0 14.6 12.4 ALP-07106F	 21.9 12.8 17.7 16.5 ALP-07206	 22.5 12.4 20.0 13.8 ALP-07306
 23.0 23.0 17.5 14.0 ALP-07406-A	 17.6 14.0 ALP-07406-O	 23.0 24.5 17.5 14.0 ALP-07406-Q	 21.4 18.0 14.2 14.3 ALP-09019/28
 9.6 11.2 16.7 8.3 14.7 ALP-09019/29	 18.9 13.2 12.5 7.1 15.6 ALP-09019/36	 22.5 19.5 15.5 15.5 ALP-09049/03	 17.6 11.6 16.6 14.4 ALP-09065/05
 19.2 24.5 16.7 ALP-09065/13	 24.1 13.5 ALP-09068/9A	 29.3 16.0 ALP-09068/9B	 17.0 11.2 14.8 ALP-09069/9
 24.3 19.3 13.3 ALP-09072/4	 17.4 18.3 ALP-09072/17	 18.5 16.4 ALP-09072/18A	 19.2 14.0 ALP-09072/18B



CELLULAR PROFILES | SPONGES

ALP-09072/24A	ALP-09072/24B	ALP-09072/27	ALP-09072/28A
ALP-09072/28B	ALP-09073/02	ALP-09073/03	ALP-09073/05
ALP-09077/10	ALP-09077/09	ALP-09083/4A	ALP-09083/4B
ALP-09083/4C	ALP-09083/5A	ALP-09083/5B	ALP-09085/5
ALP-09085/9	ALP-09085/11	ALP-09085/22	ALP-09085/23
ALP-09085/24	ALP-09087/65	ALP-09088/1	ALP-09088/2A



CELLULAR PROFILES | SPONGES

ALP-09088/2B	ALP-09088/3	ALP-09088/4	ALP-11012
ALP-11025	ALP-12106	ALP-12605	ALP-12908
ALP-14006	ALP-20006	ALP-09035/11	ALP-09068/06
ALP-09072/01	ALP-09076/06	ALP-09084/01A	ALP-09084/01B
ALP-09084/03A	ALP-09084/03B	ALP-09087/30	ALP-09087/31
ALP-09087/33	ALP-09087/34	ALP-09087/35	ALP-09087/36

ALP-11012

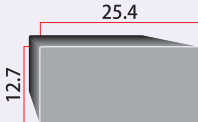
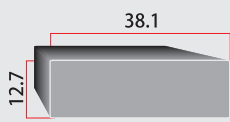
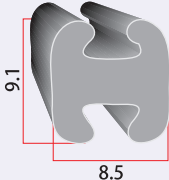
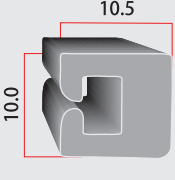
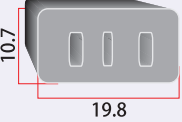
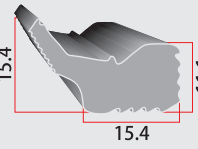
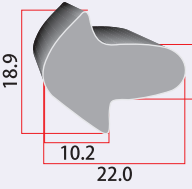
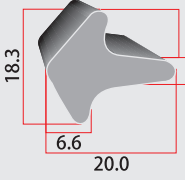
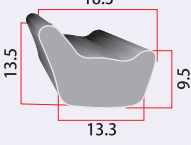
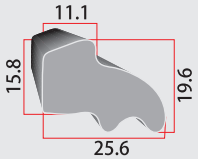
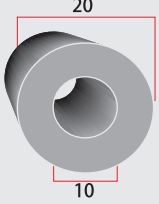
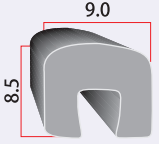
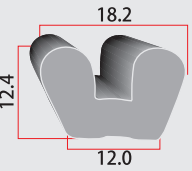
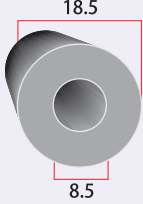
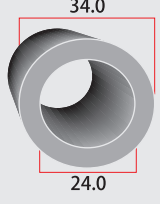
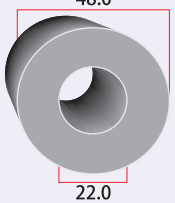
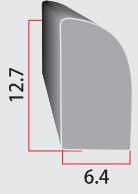
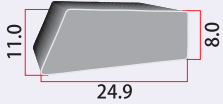
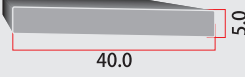
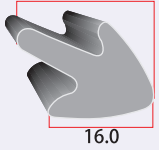
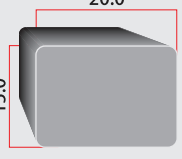
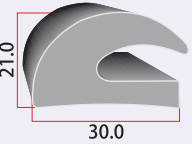
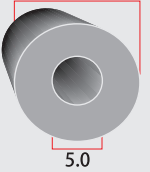


CELLULAR PROFILES | SPONGES

ALP-09087/37	ALP-09089/1	ALP-09089/2	ALP-09087/3
ALP-09087/4	ALP-09087/7	ALP-09087/10	ALP-09069/22
ALP-09072/1	ALP-09072/3	ALP-09019/35	ALP-09019/12
ALP-09019/13	ALP-09019/14	ALP-09019/15	ALP-09019/16
ALP-09019/17	ALP-09019/24	ALP-09019/25	ALP-09019/26
ALP-09019/37	ALP-09019/38	ALP-09035/07	ALP-09035/08



CELLULAR PROFILES | SPONGES

 25.4 12.7 ALP-09035/09	 38.1 12.7 ALP-09035/10	 9.1 8.5 ALP-07405	 10.5 10.0 ALP-07407
 10.7 19.8 ALP-09019/27	 15.4 11.1 15.4 ALP-09019/30	 18.9 10.2 22.0 8.6 ALP-09019/31	 18.3 7.9 3.5 6.6 20.0 ALP-09019/32
 18.3 13.5 9.5 13.3 ALP-09019/33	 11.1 15.8 19.6 25.6 ALP-09019/34	 6.5 24.8 ALP-19006	 20 10 ALP-19034
 9.0 8.5 ALP-19109	 18.2 12.4 12.0 ALP-19110	 18.5 8.5 ALP-19071	 34.0 24.0 ALP-19074
 48.0 22.0 ALP-19075	 12.7 6.4 ALP-19065	 11.0 24.9 8.0 ALP-07208	 5.0 40.0 ALP-09024/21
 23.0 16.0 ALP-09033/8	 20.0 15.0 ALP-09035/12	 21.0 30.0 ALP-09087/45	 11.0 5.0 ALP-19164

Master

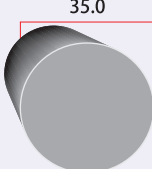
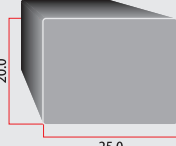
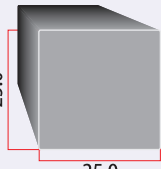
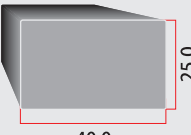
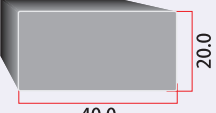
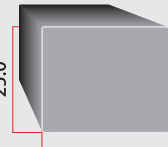
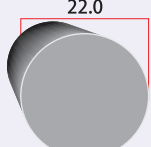
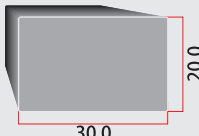
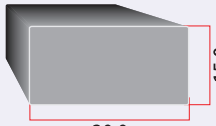
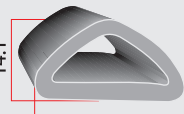
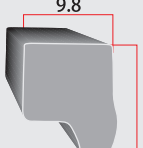
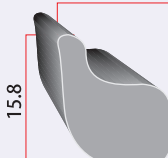
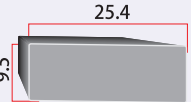
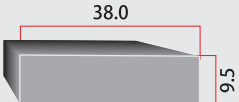
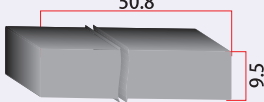
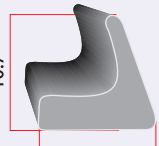
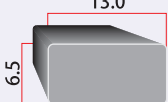
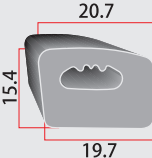
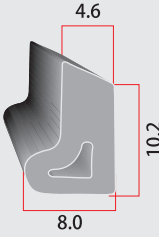


CELLULAR PROFILES | SPONGES

ALP-19165	ALP-19166	ALP-19167	ALP-19168
ALP-19173	ALP-19174	ALP-19175	ALP-19176
ALP-19177	ALP-19178	ALP-09042/20	ALP-09042/21
ALP-09042/10	ALP-19086	ALP-19091	ALP-09042/14
ALP-09042/15	09GC089	09GC106	09GE180
09GE182	ALP-09051/8	ALP-09061/1	ALP-09087/20



CELLULAR PROFILES | SPONGES

 35.0 ALP-09087/21	 20.0 25.0 ALP-09087/22	 25.0 25.0 ALP-09087/23	 40.0 25.0 ALP-09087/24
 40.0 20.0 ALP-09087/25	 25.0 30.0 ALP-09087/26	 22.0 ALP-09087/27	 30.0 20.0 ALP-09087/28
 30.0 15.0 ALP-09087/29	 14.1 26.5 ALP-15005	 3.0 15.7 13.6 ALP-07014	 9.8 13.0 ALP-11013
 13.3 15.8 ALP-21004	 19.0 9.5 ALP-09042/16	 25.4 9.5 ALP-09042/17	 38.0 9.5 ALP-09042/18
 50.8 9.5 ALP-09042/19	 13.7 9.8 ALP-09027/1	 10.7 10.6 ALP-29008	 20.0 32.0 ALP-09087/43
 13.0 6.5 ALP-09087/44	 20.7 15.4 19.7 ALP-09084/06	 15.2 4.7 09GE068	 4.6 10.2 8.0 ALP-09072/5



CELLULAR PROFILES | SPONGES

ALP - 09AD63	ALP - 09AD51	ALP - 09AD55	ALP - 09AD64
ALP - 09AD52	ALP - 09AD67	ALP - 09AD65	ALP - 09AD53
ALP - 09AD69	ALP - 09AD71	ALP - 07ES16	ALP - 09AD70
ALP - 07ES13	ALP - 070M14	ALP - 09AD75	ALP - 07ES06RR
ALP - 07YE06F A-A	ALP - 07YE16	ALP - 070M09	ALP - 07YE06F B-B
ALP - 031508	09GE095	ALP - 09FC01	ALP - 031509 A-A



CELLULAR PROFILES | SPONGES

ALP - 09FC04	ALP - 09FC03	09GE071	ALP - 07ES05
ALP - 09AD01	ALP - 09AD09	ALP - 09AD13	ALP - 09AD02
ALP - 09AD10	ALP - 09AD74	ALP - 09AD03	ALP - 09AD11
ALP - 09AD77	ALP - 09AD04	ALP - 09AD12	ALP - 09AD101
ALP - 09AD76	ALP - 07OM09	ALP - 07YE06R B-B	ALP - 07YE06R Q-Q
ALP - 03IT09 A-A	ALP - 03IT15	ALP - 03IT06RR B-B	ALP - 03IT09 D-D

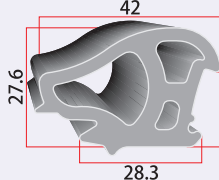
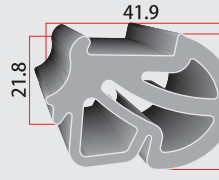
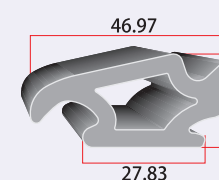
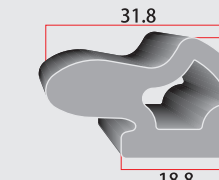
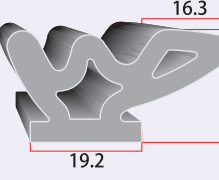
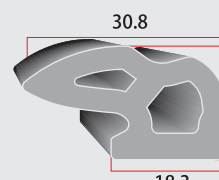
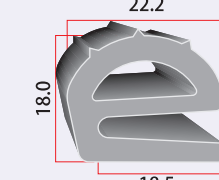
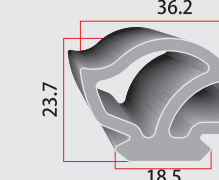
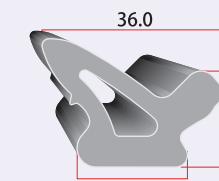
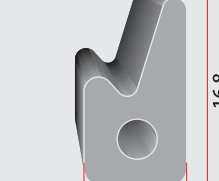
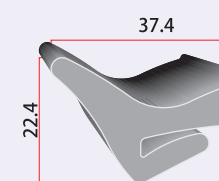
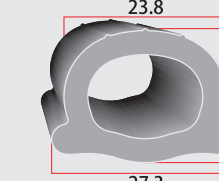
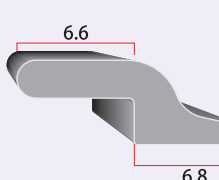
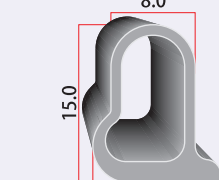
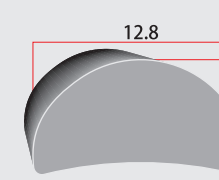
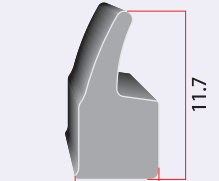
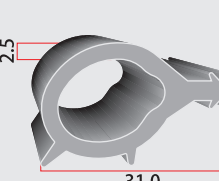
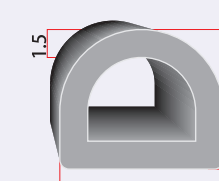
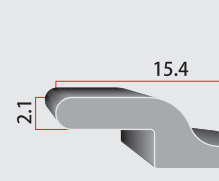
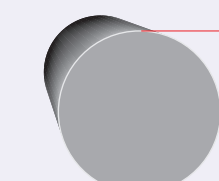
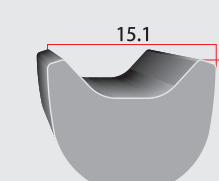
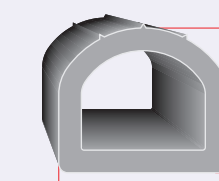
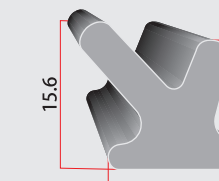


CELLULAR PROFILES | SPONGES

ALP - 03IT14	ALP - 03IT06RR E-E	ALP - 03IT08	ALP - 03IT13
ALP - 078C06F A-A	ALP - 03IT05	ALP - 03IT06RR A-A	ALP - 07GP06
ALP - 07GP16	ALP - 09M103	ALP - 09M101	ALP - 09AD102
01ML02	09MI03	09MI01	09FC02
12IN08	07OM14	09GE099	09GE131
09GE135	09GE136	09GE137	09GE142



CELLULAR PROFILES | SPONGES

 09GA017A	 09GA017B	 09GA109	 09GA206A
 09GA206B	 09GA206C	 09067/10A	 09067/10B
 ALP 09GA203	 09GE069	 09GB131	 09GB134
 09GC050	 09GC057	 09GC082	 09GC088
 09GC105	 09GC108	 09GC109	 09GC114
 09GC117	 09GC118	 09GC119	 09GC136

Meister



CELLULAR PROFILES | SPONGES

09GC137	09GC138	09GC139	09GC140
09GC141	09GC142	09GC143	09GC144
09GC145	09GD120	09GE004	09GE040
09GE056	09GE062		



CELLULAR PROFILES WITH NYLON INSERTS

07YE06R Q-Q	07YE06F A-A	07MM06 CC	07YE06F B-B
07MM06 BB	07ME08	07YE06R B-B'	07MM06 AA
07MK06 AA	07MK06 BB	07MM06 BC	07MB06 A-A
07MB06 B-B	03ITO6RR B-B	01ML06	07MA06
03ITO6RR E-E	01ML06 B-B	078C06F A-A	078C06 F-F
03ITO6RR A-A	03I508	03IT09 A-A	03I509 A-A



CELLULAR PROFILES WITH NYLON INSERTS

01ML05	03I509 B-B	05NI06AA	08MW06BB
18TB05	05NI06BB	12MP15	12605
05NI05	7010	08MW06AA	12NV08

