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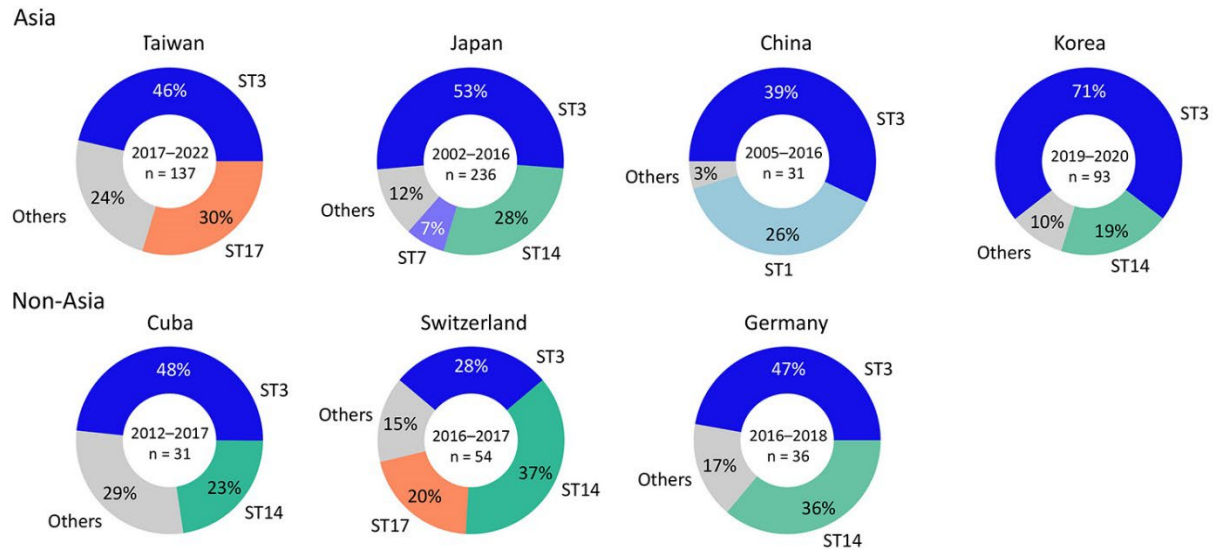
# Macrolide-Resistant *Mycoplasma pneumoniae* Infections among Children before and during COVID-19 Pandemic, Taiwan, 2017–2023

## Appendix

**Appendix Table.** Relationship between year of isolation and sequence type (ST) among *M. pneumoniae* isolates, 2017–2023

Year	ST3	ST17	ST9	ST20	ST33	Others*	Nontypable	Total
2017	12	14	3	0	2	8	9	48
2018	24	14	1	0	2	7	10	58
2019	18	7	0	0	0	5	2	32
2020	0	3	0	0	0	1	17	21
2021	6	2	0	0	0	0	3	11
2022	3	0	0	3	0	0	9	15
2023	17	0	0	1	0	2	4	24
Total	80	40	4	4	4	23	54	209

\*Other STs include ST5, ST12, ST26, ST30, ST36, ST39, and ST46.



**Appendix Figure.** The distribution of *M. pneumoniae* strains from Japan (1), China (2), South Korea (3), Cuba (4), Switzerland (5), Germany (5), and Taiwan (2), 2002–2022. ST3 was the predominant strain in these countries, while other strains vary from one country to another. ST, sequence type.

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