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| --- | --- | --- | --- | --- | --- | --- | --- |
| Study | n | Env\_matter | Study Centre | Country | Target region | Sequence platform | Qiita study ID |
| Obese\_twins | 281 | feces | CGS-GL | US | V2 | 454 GS FLX | 77 |
| Malawian\_twins\_V2 | 1042 | feces | CGS\_GL | Malawi, US | unknown | 454 FLX | 737 |
| Atherosclerosis | 73 | feces | Cornell University | Sweden | V2 | 454 FLX | 349 |
| Space\_time | 602 | feces | CCME | US | V2 | 454 FLX | 449 |
| American\_gut | 563 | feces | UCSDMI | US, Canada, UK, Australia, Sweden, Norway, New Zealand | V4 | Illumina | 10317 |
| Culture\_age\_family | 528 | feces |  | Venezuela, Malawi, US | V4 | Illumina | 850 |
| Storage\_conditions | 144 | feces | CCME | US | V2 | 454 FLX | 353 |
| Dietary\_enterotype | 95 | feces | UPENN | US | V2 | 454 FLX | 1010 |
| Fermilab\_spatial | 708 | soil | CCME | US | V4 | Illumina | 990 |
| Canadian\_metamicrobiome | 13 | soil | CCME | Canada | V4 | Illumina | 632 |
| Neon\_soils | 341 | soil | CCME | US | V4 | 454 | 619 |
| pH\_metagenome | 14 | soil | CCME | Scotland | V4 | Illumina | 805 |