```
In [1]: import requests
        import pandas as pd
        import matplotlib.pyplot as plt
        # this is the server which hosts all the patient data.
        # in the real-world this would be an EHR vendor
        # this is a open-source server with synthetic data for developers to learn a
        FHIR BASE = "http://hapi.fhir.org/baseR4"
In [2]: # Healthcare applications rarely store all their own data. Instead, they
        # exchange information with electronic health record (EHR) systems using
        # standards. FHIR (Fast Healthcare Interoperability Resources) is a standard
        # based on RESTful web API ideas and allows for retrieving patient informati
        # from an EHR AND adding information to an EHR (provided the application has
        # authorization)
        # In this tutorial, we will walk through how to interact with a public FHIR
        # server to get information about a patient.
        # We will also create a synthetic patient with clinical information (because
        # public servers often don't have the test patients we might need)
        # Pretend you are building a new health application for patients which will
        # them some health trends and their current medications. The app should be a
        # to:

    Retrieve health data such as blood pressure, heart rate, and lab resul

            - Retrieve active medications
        # Pull this information back out and display it as tables and charts so pati
        # can see trends in their health over time
        # By the end of this exercise, you'll have a toy prototype that demonstrates
        # a modern health application can connect to a standards-based health data
        # source.
In [3]: # let's query the FHIR server to get information about a patient.
        # we'll find up to 5 patients with last name "Smith" (chosen because these
        # public servers often use this name)
        # Recall that everything in FHIR is a "Resource" (patients, observations, et
        resp = requests.get(f"{FHIR_BASE}/Patient", params={"name": "Smith", "_count
        patients = resp.json()
        for entry in patients.get("entry", []):
            patient = entry["resource"]
            print(patient["id"], patient.get("name"))
```

```
597226 [{'use': 'official', 'family': 'Smith', 'given': ['Sean']}]
597372 [{'family': 'Joe', 'given': ['Smith']}]
597502 [{'family': 'Joe', 'given': ['Smith']}]
597509 [{'family': 'Joe', 'given': ['Smith']}]
618626 [{'use': 'official', 'text': 'Julia Smith', 'family': 'Smith', 'given': ['Julia']}]

In [41: # we can see that there are 3 patients with this last name. each has a uniqu# let's look at "Observations" from the first, limiting to the first 50

patient_id = "pat2" # replace with real ID from above
resp = requests.get(f"{FHIR_BASE}/Observation", params={"patient": patient_i obs = resp.json()

# and now let's print out all the observations
obs

# it's a lot! copy into a JSON viewer to look at the structure
# https://jsonviewer.stack.hu/
```

```
Out[4]: {'resourceType': 'Bundle',
          'id': '5ce8ae1d-b358-4f20-813c-710fe72ec8f0',
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         10fe72ec8f0& getpagesoffset=50& count=50& pretty=true& bundletype=searchse
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y Automated count'}],
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t'}],
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<b>code</b>: Alanine aminotransferase [Enzymatic activity/volume] in
Serum or Plasma <span style="background: LightGoldenRodYellow; margin: 4px;
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a href="https://browser.ihtsdotools.org/">SNOMED CT</a>#34608000)</span></p
><b>subject</b>: <a href="patient-example-b.html">Patient/pat2</a> &quot
;Duck DONALD"<b>performer</b>: <a href="organization-example-la">-example-la</a>
b.html">0rganization/1832473e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory,
Inc</a> &quot;Clinical Lab&quot;<b>value</b>: 54 U/L<b>interp
retation</b>: High <span style="background: LightGoldenRodYellow; margin: 4
px; border: 1px solid khaki"> (<a href="http://terminology.hl7.org/5.1.0/Co
deSystem-v3-ObservationInterpretation.html">ObservationInterpretation</a>#
H)</span><h3>ReferenceRanges</h3>-+
<b>Low</b>+td>45
U/L</div>'},
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Serum or Plasma'},
     {'system': 'http://snomed.info/sct', 'code': '34608000'}]},
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   'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
     'display': 'Acme Laboratory, Inc'}],
   'valueQuantity': {'value': 54, 'unit': 'U/L'},
    'interpretation': [{'coding': [{'system': 'http://terminology.hl7.org/C
odeSystem/v3-ObservationInterpretation',
       'code': 'H'}]}],
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     'high': {'value': 45, 'unit': 'U/L'}}]},
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    'text': {'status': 'generated',
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ative: Observation</b><a name="o15"> </a><div style="display: inline-bl
ock; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px soli
d #8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0
px">Resource Observation "o15" </div><b>status</b>: final
<b>code</b>: Aspartate aminotransferase [Enzymatic activity/volume]
in Serum or Plasma <span style="background: LightGoldenRodYellow; margin: 4
px; border: 1px solid khaki"> (<a href="https://loinc.org/">LOINC</a>#192
0-8; <a href="https://browser.ihtsdotools.org/">SNOMED CT</a>#45896001)</sp
an><b>subject</b>: <a href="patient-example-b.html">Patient/pat2</a>
"Duck DONALD"<b>performer</b>: <a href="organization-examp"
le-lab.html">0rganization/1832473e-2fe0-452d-abe9-3cdb9879522f: Acme Labora
tory, Inc</a> &quot;Clinical Lab&quot;<b>value</b>: 30 U/L<h3>Re
ferenceRanges</h3>->b>Low</b>
<b>High</b>*</table
e></div>'},
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   'code': {'coding': [{'system': 'http://loinc.org',
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      'display': 'Aspartate aminotransferase [Enzymatic activity/volume] i
n Serum or Plasma'},
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{'system': 'http://snomed.info/sct', 'code': '45896001'}]},
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879522f',
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     'high': {'value': 41, 'unit': 'U/L'}}]},
   'search': {'mode': 'match'}},
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    'id': '44801538',
    'meta': {'versionId': '1',
    'lastUpdated': '2024-07-11T15:44:08.135+00:00',
    'source': '#tYPNjWj0MUsuCAKv'},
    'text': {'status': 'generated',
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ative: Observation</b><a name="o6"> </a><div style="display: inline-blo
ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
">Resource Observation "o6" </div><b>status</b>: final
<b>code</b>: Glucose [Moles/volume] in Serum or Plasma <span style="back"
ground: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a hr
ef="https://loinc.org/">LOINC</a>#14749-6; <a href="https://browser.ihtsdot
ools.org/">SNOMED CT</a>#36048009)</span><b>subject</b>: <a href="pa
tient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b>per
former</b>: <a href="organization-example-lab.html">Organization/1832473e-2
fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab&quo
t;<b>value</b>: 7.4 mmol/L<h3>ReferenceRanges</h3><table class
="grid">-b>Low</b>b>High</b>*
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    'subject': {'reference': 'Patient/pat2'},
    'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
     'display': 'Acme Laboratory, Inc'}],
    'valueQuantity': {'value': 7.4, 'unit': 'mmol/L'},
    'referenceRange': [{'low': {'value': 3, 'unit': 'mmol/L'},
     'high': {'value': 7.7, 'unit': 'mmol/L'}}]},
   'search': {'mode': 'match'}},
  {'fullUrl': 'https://hapi.fhir.org/baseR4/Observation/44801539',
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    'id': '44801539',
    'meta': {'versionId': '1',
    'lastUpdated': '2024-07-11T15:44:08.135+00:00',
    'source': '#tYPNjWj0MUsuCAKv'},
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ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
">Resource Observation "o7" </div><b>status</b>: final
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```
<b>code</b>: Urea nitrogen [Moles/volume] in Serum or Plasma <span style
="background: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki">
(<a href="https://loinc.org/">LOINC</a>#14937-7; <a href="https://browser.i")</pre>
htsdotools.org/">SNOMED CT</a>#273967009)</span><b>subject</b>: <a h
ref="patient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<p
><b>performer</b>: <a href="organization-example-lab.html">0rganization/183
2473e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical
Lab"<b>value</b>: 4.7 mmol/L<h3>ReferenceRanges</h3><table
class = "qrid" >   -  < tb > Low < /b >                                                                         
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     {'system': 'http://snomed.info/sct', 'code': '273967009'}]},
    'subject': {'reference': 'Patient/pat2'},
    'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
     'display': 'Acme Laboratory, Inc'}],
    'valueQuantity': {'value': 4.7, 'unit': 'mmol/L'},
    'referenceRange': [{'low': {'value': 2, 'unit': 'mmol/L'},
     'high': {'value': 7, 'unit': 'mmol/L'}}]},
   'search': {'mode': 'match'}},
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    'source': '#tYPNjWj0MUsuCAKv'},
    'text': {'status': 'generated',
    'div': '<div xmlns="http://www.w3.org/1999/xhtml"><b>Generated Narr
ative: Observation</b><a name="o8"> </a><div style="display: inline-blo
ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
">Resource Observation "o8" </div><b>status</b>: final
<b>code</b>: Creatinine [Moles/volume] in Serum or Plasma <span style="b
ackground: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a
href="https://loinc.org/">LOINC</a>#14682-9; <a href="https://browser.ihtsd
otools.org/">SNOMED CT</a>#70901006)</span><b>subject</b>: <a href
="patient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b
>performer</b>: <a href="organization-example-lab.html">0rganization/183247
3e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab
"<b>value</b>: 0.09 mmol/L<h3>ReferenceRanges</h3><table cl
ass="grid">-b>Low</b>b>High</b><t
d>*0.04 mmol/L0.11 mmol/L</div>'},
    'status': 'final',
    'code': {'coding': [{'system': 'http://loinc.org',
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    'subject': {'reference': 'Patient/pat2'},
    'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
      'display': 'Acme Laboratory, Inc'}],
    'valueQuantity': {'value': 0.09, 'unit': 'mmol/L'},
    'referenceRange': [{'low': {'value': 0.04, 'unit': 'mmol/L'},
```

```
'high': {'value': 0.11, 'unit': 'mmol/L'}}]},
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    'text': {'status': 'generated',
    'div': '<div xmlns="http://www.w3.org/1999/xhtml"><b>Generated Narr
ative: Observation</b><a name="o17"> </a><div style="display: inline-bl
ock; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px soli
d #8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0
px">Resource Observation "o17" </div><b>status</b>: final
<b>code</b>: Calcium [Moles/volume] in Serum or Plasma <span style="
background: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<
a href="https://loinc.org/">LOINC</a>#2000-8; <a href="https://browser.ihts"
dotools.org/">SNOMED CT</a>#71878006)</span><bssubject</b>: <a href
="patient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b
>performer</b>: <a href="organization-example-lab.html">0rganization/183247
3e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab
"<b>value</b>: 2.38 mmol/L<h3>ReferenceRanges</h3><table cl
ass="grid">-b>Low</b>b>High</b><t
d>*2.25 mmol/L2.65 mmol/L</div>'},
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879522f',
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    'valueQuantity': {'value': 2.38, 'unit': 'mmol/L'},
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   'search': {'mode': 'match'}},
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    'meta': {'versionId': '1',
    'lastUpdated': '2024-07-11T15:44:08.135+00:00',
    'source': '#tYPNjWj0MUsuCAKv'},
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ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
">Resource Observation "o1" </div><b>status</b>: final
<b>code</b>: Sodium [Moles/volume] in Serum or Plasma <span style="backg"
round: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a hre
f="https://loinc.org/">L0INC</a>#2951-2; <a href="https://browser.ihtsdotoo
ls.org/">SNOMED CT</a>#104934005)</span><b>subject</b>: <a href="pat
ient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b>perf
ormer</b>: <a href="organization-example-lab.html">0rganization/1832473e-2f
e0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab&quo
```

```
t;<b>value</b>: 140 mmol/L<h3>ReferenceRanges</h3><table class
="grid">-<b>Low</b>>b>High</b>*
137 mmol/L147 mmol/L</div>'},
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879522f',
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         'high': {'value': 147, 'unit': 'mmol/L'}}]},
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ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
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ckground: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a</pre>
href="https://loinc.org/">LOINC</a>#2823-3; <a href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://browser.ihtsdo">href="https://
tools.org/">SNOMED CT</a>#59573005)<math></span><b>subject<math></b>: <a href="p
atient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b>pe
rformer</b>: <a href="organization-example-lab.html">Organization/1832473
e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab&
quot;<b>value</b>: 4.2 mmol/L<h3>ReferenceRanges</h3>
s="qrid">-b>Low</b>High</b>= "qrid">-+ 10 -
*3.5 mmol/L5.0 mmol/L</div>'},
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879522f',
          'display': 'Acme Laboratory, Inc'}],
       'valueQuantity': {'value': 4.2, 'unit': 'mmol/L'},
       'referenceRange': [{'low': {'value': 3.5, 'unit': 'mmol/L'},
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       'id': '44801544',
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<b>code</b>: Chloride [Moles/volume] in Serum or Plasma <span style="bac"
kground: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a h
ref="https://loinc.org/">LOINC</a>#2075-0; <a href="https://browser.ihtsdot
ools.org/">SNOMED CT</a>#46511006)</span><b>subject</b>: <a href="pa
tient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b>per
former</b>: <a href="organization-example-lab.html">0rganization/1832473e-2
fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab&quo
t;<b>value</b>: 105 mmol/L<h3>ReferenceRanges</h3><table class
="grid">-<b>Low</b>>b>High</b>*
96 mmol/L109 mmol/L</div>'},
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879522f',
     'display': 'Acme Laboratory, Inc'}],
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ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
">Resource Observation "o4" </div><b>status</b>: final
<b>code</b>: Bicarbonate [Moles/volume] in Serum <span style="background"
: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a href="ht
tps://loinc.org/">LOINC</a>#1963-8; <a href="https://browser.ihtsdotools.or
g/">SNOMED CT</a>#88645003)</span><b>subject</b>: <a href="patient-e">
xample-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b>performer
</b>: <a href="organization-example-lab.html">0rganization/1832473e-2fe0-45
2d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab&quot;
<b>value</b>: 26 mmol/L<h3>ReferenceRanges</h3><
25 mmol/L33 mmol/L</div>'},
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```

```
{'system': 'http://snomed.info/sct', 'code': '88645003'}]},
    'subject': {'reference': 'Patient/pat2'},
    'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
      'display': 'Acme Laboratory, Inc'}],
    'valueQuantity': {'value': 26, 'unit': 'mmol/L'},
    'referenceRange': [{'low': {'value': 25, 'unit': 'mmol/L'},
     'high': {'value': 33, 'unit': 'mmol/L'}}]},
   'search': {'mode': 'match'}},
  {'fullUrl': 'https://hapi.fhir.org/baseR4/Observation/44801546',
   'resource': {'resourceType': 'Observation',
    'id': '44801546',
    'meta': {'versionId': '1',
    'lastUpdated': '2024-07-11T15:51:37.783+00:00',
    'source': '#kZ3KAReIzyBIPCpk'},
    'text': {'status': 'generated',
     'div': '<div xmlns="http://www.w3.org/1999/xhtml"><b>Generated Narr
ative: Observation</b><a name="o14"> </a><div style="display: inline-bl
ock; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px soli
d #8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0
px">Resource Observation "o14" </div><b>status</b>: final
<b>code</b>: Alanine aminotransferase [Enzymatic activity/volume] in
Serum or Plasma <span style="background: LightGoldenRodYellow; margin: 4px;
border: 1px solid khaki"> (<a href="https://loinc.org/">LOINC</a>#1742-6; <
a href="https://browser.ihtsdotools.org/">SNOMED CT</a>#34608000)</span></p
><b>subject</b>: <a href="patient-example-b.html">Patient/pat2</a> &guot
;Duck DONALD"<b>performer</b>: <a href="organization-example-la">- la</a>
b.html">0rganization/1832473e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory,
Inc</a> &quot;Clinical Lab&quot;<b>value</b>: 54 U/L<b>interp
retation</b>: High <span style="background: LightGoldenRodYellow; margin: 4
px; border: 1px solid khaki"> (<a href="http://terminology.hl7.org/5.1.0/Co
deSystem-v3-ObservationInterpretation.html">ObservationInterpretation</a>#
H)</span><h3>ReferenceRanges</h3>-
U/L</div>'},
    'status': 'final',
    'code': {'coding': [{'system': 'http://loinc.org',
      'code': '1742-6',
      'display': 'Alanine aminotransferase [Enzymatic activity/volume] in
Serum or Plasma'},
     {'system': 'http://snomed.info/sct', 'code': '34608000'}]},
    'subject': {'reference': 'Patient/pat2'},
    'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
     'display': 'Acme Laboratory, Inc'}],
    'valueQuantity': {'value': 54, 'unit': 'U/L'},
    'interpretation': [{'coding': [{'system': 'http://terminology.hl7.org/C
odeSystem/v3-ObservationInterpretation',
       'code': 'H'}]}],
    'referenceRange': [{'low': {'value': 0, 'unit': 'U/L'},
     'high': {'value': 45, 'unit': 'U/L'}}]},
   'search': {'mode': 'match'}},
  {'fullUrl': 'https://hapi.fhir.org/baseR4/Observation/44801547',
   'resource': {'resourceType': 'Observation',
    'id': '44801547',
    'meta': {'versionId': '1',
```

```
'lastUpdated': '2024-07-11T15:51:37.783+00:00',
    'source': '#kZ3KAReIzyBIPCpk'},
    'text': {'status': 'generated',
    'div': '<div xmlns="http://www.w3.org/1999/xhtml"><b>Generated Narr
ative: Observation</b><a name="o15"> </a><div style="display: inline-bl
ock; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px soli
d #8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0
px">Resource Observation "o15" </div><b>status</b>: final
<b>code</b>: Aspartate aminotransferase [Enzymatic activity/volume]
in Serum or Plasma <span style="background: LightGoldenRodYellow; margin: 4
px; border: 1px solid khaki"> (<a href="https://loinc.org/">LOINC</a>#192
0-8; <a href="https://browser.ihtsdotools.org/">SNOMED CT</a>#45896001)</sp
an><b>subject</b>: <a href="patient-example-b.html">Patient/pat2</a>
"Duck DONALD"<b>performer</b>: <a href="organization-examp">-examp</a>
le-lab.html">Organization/1832473e-2fe0-452d-abe9-3cdb9879522f: Acme Labora
tory, Inc</a> &quot;Clinical Lab&quot;<b>value</b>: 30 U/L<h3>Re
ferenceRanges</h3>->b>Low</b>
<b>High</b>*</table
e></div>'},
   'status': 'final',
    'code': {'coding': [{'system': 'http://loinc.org',
      'code': '1920-8',
      'display': 'Aspartate aminotransferase [Enzymatic activity/volume] i
n Serum or Plasma'},
     {'system': 'http://snomed.info/sct', 'code': '45896001'}]},
    'subject': {'reference': 'Patient/pat2'},
    'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
879522f',
     'display': 'Acme Laboratory, Inc'}],
    'valueQuantity': {'value': 30, 'unit': 'U/L'},
   'referenceRange': [{'low': {'value': 0, 'unit': 'U/L'},
     'high': {'value': 41, 'unit': 'U/L'}}]},
   'search': {'mode': 'match'}},
  {'fullUrl': 'https://hapi.fhir.org/baseR4/0bservation/44801548',
   'resource': {'resourceType': 'Observation',
   'id': '44801548',
   'meta': {'versionId': '1',
    'lastUpdated': '2024-07-11T15:51:37.783+00:00',
    'source': '#kZ3KAReIzyBIPCpk'},
   'text': {'status': 'generated',
    'div': '<div xmlns="http://www.w3.org/1999/xhtml"><b>Generated Narr
ative: Observation</b><a name="o6"> </a><div style="display: inline-blo
ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
#8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
">Resource Observation "o6" </div><b>status</b>: final
<b>code</b>: Glucose [Moles/volume] in Serum or Plasma <span style="back"
ground: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki"> (<a hr
ef="https://loinc.org/">LOINC</a>#14749-6; <a href="https://browser.ihtsdot
ools.org/">SNOMED CT</a>#36048009)</span><b>subject</b>: <a href="pa
tient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<b>per
former</b>: <a href="organization-example-lab.html">0rganization/1832473e-2
fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical Lab&quo
t;<b>value</b>: 7.4 mmol/L<h3>ReferenceRanges</h3><table class
="grid">-<b>Low</b>>b>High</b>*
'status': 'final',
```

```
'code': '14749-6',
               'display': 'Glucose [Moles/volume] in Serum or Plasma'},
              {'system': 'http://snomed.info/sct', 'code': '36048009'}]},
            'subject': {'reference': 'Patient/pat2'},
            'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
        879522f',
              'display': 'Acme Laboratory, Inc'}],
            'valueQuantity': {'value': 7.4, 'unit': 'mmol/L'},
            'referenceRange': [{'low': {'value': 3, 'unit': 'mmol/L'},
              'high': {'value': 7.7, 'unit': 'mmol/L'}}]},
           'search': {'mode': 'match'}},
          {'fullUrl': 'https://hapi.fhir.org/baseR4/0bservation/44801549',
           'resource': {'resourceType': 'Observation',
            'id': '44801549',
            'meta': {'versionId': '1',
             'lastUpdated': '2024-07-11T15:51:37.783+00:00',
             'source': '#kZ3KAReIzyBIPCpk'},
            'text': {'status': 'generated',
             'div': '<div xmlns="http://www.w3.org/1999/xhtml"><b>Generated Narr
        ative: Observation</b><a name="o7"> </a><div style="display: inline-blo
        ck; background-color: #d9e0e7; padding: 6px; margin: 4px; border: 1px solid
        #8da1b4; border-radius: 5px; line-height: 60%"><p style="margin-bottom: 0px
        ">Resource Observation "o7" </div><b>status</b>: final
        <b>code</b>: Urea nitrogen [Moles/volume] in Serum or Plasma <span style
        ="background: LightGoldenRodYellow; margin: 4px; border: 1px solid khaki">
        (<a href="https://loinc.org/">LOINC</a>#14937-7; <a href="https://browser.i")</pre>
        htsdotools.org/">SNOMED CT</a>#273967009)</span><b>subject</b>: <a h
        ref="patient-example-b.html">Patient/pat2</a> &quot;Duck DONALD&quot;<p
        ><b>performer</b>: <a href="organization-example-lab.html">Organization/183
        2473e-2fe0-452d-abe9-3cdb9879522f: Acme Laboratory, Inc</a> &quot;Clinical
        Lab"<b>value</b>: 4.7 mmol/L<h3>ReferenceRanges</h3><table
        class="grid">-<b>Low</b>>b>High</b>
        'status': 'final',
            'code': {'coding': [{'system': 'http://loinc.org',
               'code': '14937-7',
               'display': 'Urea nitrogen [Moles/volume] in Serum or Plasma'},
              {'system': 'http://snomed.info/sct', 'code': '273967009'}]},
            'subject': {'reference': 'Patient/pat2'},
            'performer': [{'reference': 'Organization/1832473e-2fe0-452d-abe9-3cdb9
        879522f',
              'display': 'Acme Laboratory, Inc'}],
            'valueQuantity': {'value': 4.7, 'unit': 'mmol/L'},
            'referenceRange': [{'low': {'value': 2, 'unit': 'mmol/L'},
              'high': {'value': 7, 'unit': 'mmol/L'}}]},
           'search': {'mode': 'match'}}]}
In [5]: # let's write some code to unpack this. the results come back as a bundle an
       # can use our json viewer to see that part of this is a list of resources
        # in each resource we can find the lab names and values!
        observations = []
        for entry in obs.get("entry", []):
           resource = entry["resource"]
```

'code': {'coding': [{'system': 'http://loinc.org',

```
if resource.get("code", {}).get("coding"):
              code = resource.get("code", {}).get("coding")[0].get("display")
              value = resource.get("valueQuantity", {}).get("value")
unit = resource.get("valueQuantity", {}).get("unit")
              time = resource.get("effectiveDateTime")
                print("Skipping entry due to missing 'coding' information.")
            if value:
                observations.append({"code": code, "value": value, "unit": unit, "ti
        df = pd.DataFrame(observations)
        print(df.head())
       Skipping entry due to missing 'coding' information.
                                                         code value
                                                                          unit time
       0
             Cholesterol [Moles/volume] in Serum or Plasma
                                                                       mmol/L None
                                                               6.3
                          Hemoglobin [Mass/volume] in Blood 176.0
       1
                                                                           g/L None
       2 Erythrocytes [#/volume] in Blood by Automated ...
                                                               5.9 x10*12/L None
       3 Hematocrit [Volume Fraction] of Blood by Autom... 55.0
                                                                            % None
       4 Erythrocyte mean corpuscular volume [Entitic v... 99.0
                                                                           fL None
In [6]: # as you can see this is a random collection of labs... for our app we need
        # detail.
        # let's create our own patient now so we have control over the data
        # we are allowed to do this because it is a public endpoint
        # this paitent will be purged at some interval
```

In [7]: # this makes a fake new patient # if everything goes well, we'll get a 201 response and an ID new patients = [{ "resourceType": "Patient", "name": [{"use": "official", "family": "Doe", "given": ["Jane"]}], "gender": "female", "birthDate": "1980-05-01" }, 'resourceType': 'Patient', 'name': [{'use': 'official', 'family': 'Smith', 'qiven': ['Martina'] 'gender': 'female', 'birthDate': '2003-12-05' } 1 for patient in new_patients: resp = requests.post(f"{FHIR_BASE}/Patient", json=patient) print(resp.status_code, resp.json()) patient id = resp.json()["id"] print("New Patient ID:", patient_id)

```
201 {'resourceType': 'Patient', 'id': '50166462', 'meta': {'versionId': '1',
       'lastUpdated': '2025-10-14T19:32:26.538+00:00', 'source': '#SLK7GQSXIsyva99
       D'}, 'text': {'status': 'generated', 'div': '<div xmlns="http://www.w3.org/1
       999/xhtml"><div class="hapiHeaderText">Jane <b>D0E </b></div><table class="h
       apiPropertyTable">Date of birthspan>01 May 1980</sp
       an></div>'}, 'name': [{'use': 'official', 'family'
       : 'Doe', 'given': ['Jane']}], 'gender': 'female', 'birthDate': '1980-05-01'}
      New Patient ID: 50166462
       201 {'resourceType': 'Patient', 'id': '50166463', 'meta': {'versionId': '1',
       'lastUpdated': '2025-10-14T19:32:27.539+00:00', 'source': '#muB8mQMNTyb8xEU
      K'}, 'text': {'status': 'generated', 'div': '<div xmlns="http://www.w3.org/1</pre>
       999/xhtml"><div class="hapiHeaderText">Martina <b>SMITH </b></div><table cla
       ss="hapiPropertyTable">Date of birth<span>05 Decembe
       r 2003</span></div>'}, 'name': [{'use': 'official'
       , 'family': 'Smith', 'given': ['Martina']}], 'gender': 'female', 'birthDate'
       : '2003-12-05'}
      New Patient ID: 50166463
In [8]: patient id = resp.json()["id"]
       print("New Patient ID:", patient_id)
      New Patient ID: 50166463
In [9]: # let's add a bunch of random data, one set of observations per day
        from random import randint, gauss
        from datetime import datetime, timedelta
        base date = datetime(2025, 8, 1)
        observations = []
        for i in range(5): # 5 days
           dt = (base_date + timedelta(days=i)).isoformat() + "Z"
           observations += [
               {
                   "resourceType": "Observation",
                   "status": "final",
                   "category": [{"coding": [{"system": "http://terminology.hl7.org/
                                            "code": "vital-signs"}]}],
                   "code": {"text": "Systolic Blood Pressure"},
                   "subject": {"reference": f"Patient/{patient id}"},
                   "effectiveDateTime": dt,
                   "valueQuantity": {"value": randint(110, 135), "unit": "mmHg"}
               },
                   "resourceType": "Observation",
                   "status": "final",
                   "category": [{"coding": [{"system": "http://terminology.hl7.org/
                                            "code": "vital-signs"}]}],
                   "code": {"text": "Diastolic Blood Pressure"},
                   "subject": {"reference": f"Patient/{patient id}"},
                   "effectiveDateTime": dt,
                   "valueQuantity": {"value": randint(70, 90), "unit": "mmHg"}
               },
                   "resourceType": "Observation",
```

```
"category": [{"coding": [{"system": "http://terminology.hl7.org/
                                                "code": "vital-signs"}]}],
                      "code": {"text": "Heart rate"},
                      "subject": {"reference": f"Patient/{patient_id}"},
                      "effectiveDateTime": dt,
                      "valueQuantity": {"value": randint(65, 90), "unit": "beats/min"}
                 },
                      "resourceType": "Observation",
                      "status": "final",
                      "category": [{"coding": [{"system": "http://terminology.hl7.org/
                                                "code": "laboratory"}]}],
                     "code": {"text": "Glucose"},
                      "subject": {"reference": f"Patient/{patient id}"},
                     "effectiveDateTime": dt,
                      "valueQuantity": {"value": round(gauss(95, 10), 1), "unit": "mg/
                 }
             1
         # POST them
         for obs in observations:
             r = requests.post(f"{FHIR_BASE}/Observation", json=obs, params={"patient"}
             print("Obs:", r.status_code, r.json().get("id"))
        Obs: 201 50166465
        0bs: 201 50166466
        Obs: 201 50166467
        Obs: 201 50166468
        Obs: 201 50166469
        Obs: 201 50166471
        Obs: 201 50166472
        Obs: 201 50166473
        Obs: 201 50166474
        0bs: 201 50166475
        Obs: 201 50166477
        Obs: 201 50166478
        Obs: 201 50166479
        Obs: 201 50166480
        Obs: 201 50166481
        0bs: 201 50166483
        Obs: 201 50166484
        Obs: 201 50166485
        0bs: 201 50166486
        0bs: 201 50166487
In [10]: # Query observations
         obs_resp = requests.get(f"{FHIR_BASE}/Observation", params={"patient": patie
         print(len(obs_resp.json().get("entry", [])), "observations found")
         print(patient_id)
        20 observations found
        50166463
In [11]: observations = []
         for entry in obs_resp.json().get("entry", []):
             resource = entry["resource"]
```

"status": "final",

```
code = resource.get("code", {}).get("text")
value = resource.get("valueQuantity", {}).get("value")
unit = resource.get("valueQuantity", {}).get("unit")
time = resource.get("effectiveDateTime")
if value:
    observations.append({"code": code, "value": value, "unit": unit, "ti

df = pd.DataFrame(observations)
print(df)
```

```
code value
                                        unit
                                                             time
0
    Systolic Blood Pressure 131.0
                                        mmHg 2025-08-01T00:00:00Z
1
   Diastolic Blood Pressure 72.0
                                        mmHg 2025-08-01T00:00:00Z
2
                 Heart rate 75.0 beats/min 2025-08-01T00:00:00Z
                                       mg/dL 2025-08-01T00:00:00Z
3
                    Glucose 100.6
4
    Systolic Blood Pressure 110.0
                                        mmHg 2025-08-02T00:00:00Z
5
   Diastolic Blood Pressure 79.0
                                        mmHg 2025-08-02T00:00:00Z
6
                 Heart rate 78.0 beats/min 2025-08-02T00:00:00Z
                    Glucose 99.5
7
                                       mg/dL 2025-08-02T00:00:00Z
                                        mmHg 2025-08-03T00:00:00Z
8
    Systolic Blood Pressure 122.0
9
    Diastolic Blood Pressure 84.0
                                        mmHg 2025-08-03T00:00:00Z
                             71.0 beats/min 2025-08-03T00:00:00Z
10
                 Heart rate
11
                    Glucose 84.3
                                       mg/dL 2025-08-03T00:00:00Z
    Systolic Blood Pressure 111.0
12
                                        mmHg 2025-08-04T00:00:00Z
   Diastolic Blood Pressure 88.0
                                        mmHg 2025-08-04T00:00:00Z
13
14
                 Heart rate 89.0 beats/min 2025-08-04T00:00:00Z
                                       mg/dL 2025-08-04T00:00:00Z
15
                    Glucose 110.0
    Systolic Blood Pressure 116.0
16
                                        mmHg 2025-08-05T00:00:00Z
17
   Diastolic Blood Pressure 90.0
                                        mmHq 2025-08-05T00:00:00Z
18
                 Heart rate 83.0 beats/min 2025-08-05T00:00:00Z
19
                    Glucose 89.0
                                       mg/dL 2025-08-05T00:00:00Z
```

```
In [12]: medications = [
                 "resourceType": "MedicationStatement",
                 "status": "active",
                 "medicationCodeableConcept": {"text": "Lisinopril 10mg tablet"},
                 "subject": {"reference": f"Patient/{patient_id}"},
                 "effectiveDateTime": "2025-09-10T11:00:00Z"
             },
                 "resourceType": "MedicationStatement",
                 "status": "active",
                 "medicationCodeableConcept": {"text": "Lisinopril 10mg tablet"},
                 "subject": {"reference": f"Patient/{patient_id}"},
                 "effectiveDateTime": "2025-09-11T11:30:00Z"
             },
             {
                 "resourceType": "MedicationStatement",
                 "status": "active",
                 "medicationCodeableConcept": {"text": "Lisinopril 20mg tablet"},
                 "subject": {"reference": f"Patient/{patient_id}"},
                 "effectiveDateTime": "2025-09-12T12:00:00Z"
             }
         ]
```

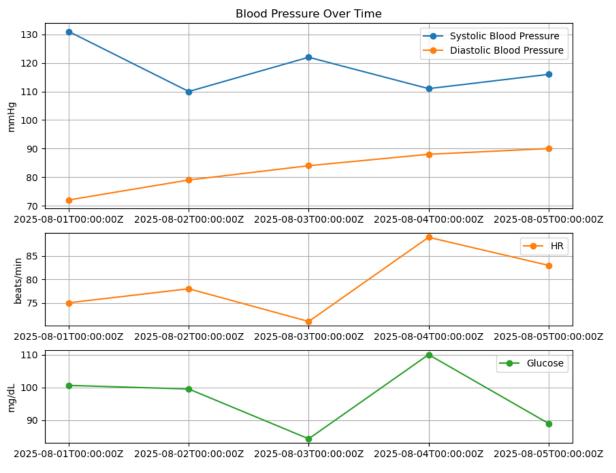
```
for med in medications:
            r = requests.post(f"{FHIR_BASE}/MedicationStatement", json=med)
            print("Med:", r.status code, r.json().get("id"))
       Med: 201 50166489
       Med: 201 50166490
       Med: 201 50166491
In [13]: patient_id
Out[13]: '50166463'
In [14]: # Now that we have our fake patient set up, let's go back to the goal and pu
        # their data
        # Observations
        obs_resp = requests.get(f"{FHIR_BASE}/Observation", params={"patient": patie
        print("----")
        print(len(obs_resp.json().get("entry", [])), "observations")
        observations = []
        for entry in obs_resp.json().get("entry", []):
            resource = entry["resource"]
            code = resource.get("code", {}).get("text")
            value = resource.get("valueQuantity", {}).get("value")
            unit = resource.get("valueQuantity", {}).get("unit")
            time = resource.get("effectiveDateTime")
            if value:
                observations.append({"code": code, "value": value, "unit": unit, "ti
        df = pd.DataFrame(observations)
        print(df)
        # Medications
        print("\n----")
        med resp = requests.get(f"{FHIR BASE}/MedicationStatement", params={"patient"}
        for e in med_resp.json().get("entry", []):
            print("Medication:", e["resource"].get("medicationCodeableConcept", {}).
```

```
20 observations
                              code value unit
                                                                    time
            Systolic Blood Pressure 131.0
                                               mmHg 2025-08-01T00:00:00Z
           Diastolic Blood Pressure 72.0 mmHg 2025-08-01T00:00:00Z
       1
                        Heart rate 75.0 beats/min 2025-08-01T00:00:00Z
       2
       3
                           Glucose 100.6 mg/dL 2025-08-01T00:00:00Z
           Systolic Blood Pressure 110.0
                                             mmHg 2025-08-02T00:00:00Z
       4
       5
           Diastolic Blood Pressure 79.0
                                               mmHg 2025-08-02T00:00:00Z
       6
                        Heart rate 78.0 beats/min 2025-08-02T00:00:00Z
       7
                           Glucose 99.5
                                              mg/dL 2025-08-02T00:00:00Z
       8
            Systolic Blood Pressure 122.0
                                              mmHg 2025-08-03T00:00:00Z
                                               mmHg 2025-08-03T00:00:00Z
       9
           Diastolic Blood Pressure 84.0
       10
                        Heart rate 71.0 beats/min 2025-08-03T00:00:00Z
       11
                           Glucose 84.3
                                          mg/dL 2025-08-03T00:00:00Z
       12
            Systolic Blood Pressure 111.0
                                             mmHg 2025-08-04T00:00:00Z
       13 Diastolic Blood Pressure 88.0 mmHg 2025-08-04T00:00Z
       14
                        Heart rate 89.0 beats/min 2025-08-04T00:00:00Z
       15
                           Glucose 110.0 mg/dL 2025-08-04T00:00:00Z
       16
           Systolic Blood Pressure 116.0
                                             mmHg 2025-08-05T00:00:00Z
       17 Diastolic Blood Pressure 90.0 mmHg 2025-08-05T00:002
                        Heart rate 83.0 beats/min 2025-08-05T00:00:00Z
       18
       19
                           Glucose 89.0 mg/dL 2025-08-05T00:00:00Z
       ----- MEDICATIONS -----
       Medication: Lisinopril 10mg tablet
       Medication: Lisinopril 10mg tablet
       Medication: Lisinopril 20mg tablet
In [15]: # ok now let's plot!
        fig, axes = plt.subplots(3, 1, figsize=(10, 8), gridspec_kw={"height_ratios"
        # show bp on one plot
        bp = df[df["code"].str.contains("Blood Pressure")]
        for code in ["Systolic Blood Pressure", "Diastolic Blood Pressure"]:
            subset = bp[bp["code"] == code]
            axes[0].plot(subset["time"], subset["value"], marker="o", label=code)
        axes[0].set_ylabel("mmHg")
        axes[0].set_title("Blood Pressure Over Time")
        axes[0].legend()
        axes[0].grid(True)
        hr = df[df["code"] == "Heart rate"]
        axes[1].plot(hr["time"], hr["value"], marker="o", color="tab:orange",label="
        axes[1].set_ylabel("beats/min")
        axes[1].legend()
        axes[1].grid(True)
        glucose = df[df["code"] == "Glucose"]
        axes[2].plot(glucose["time"], glucose["value"], marker="o", color="tab:green
        axes[2].set_ylabel("mg/dL")
        axes[2].legend()
        axes[2].grid(True)
```

----- OBSEVATIONS -----

IHI_assignment2





```
In [ ]: # if you made it this far congratulations! To extend your knowledge, please
    # find a new resource type on the FHIR specification, make some synthetic da
    # and retrieve it, showing the data however you like.

# you can browse resources here: https://build.fhir.org/resourcelist.html
```

Make synthetic data with immunization

```
Out[17]: {'reference': 'Patient/49233412'}
In [18]: {
           "resourceType": "Immunization",
           "status": "completed",
           "vaccineCode": {
             "coding": [
                 "system": "http://hl7.org/fhir/sid/cvx",
                 "code": "102",
                 "display": "MMR"
               }
             1
           },
           "patient": {
             "reference": "Patient/49233412"
           },
           "occurrenceDateTime": "2025-08-15T10:00:00Z",
           "performer": [
             {
               "function": {
                 "coding": [
                     "system": "http://terminology.hl7.org/CodeSystem/v2-0443",
                     "code": "AP",
                     "display": "Administering Provider"
                 1
               },
               "actor": {
                 "reference": "Practitioner/example-practitioner-id"
               }
             }
           ],
           "lotNumber": "ABC12345",
           "expirationDate": "2026-08-15"
Out[18]: {'resourceType': 'Immunization',
           'status': 'completed',
           'vaccineCode': {'coding': [{'system': 'http://hl7.org/fhir/sid/cvx',
              'code': '102',
              'display': 'MMR'}]},
           'patient': {'reference': 'Patient/49233412'},
           'occurrenceDateTime': '2025-08-15T10:00:00Z',
           'performer': [{'function': {'coding': [{'system': 'http://terminology.hl
          7.org/CodeSystem/v2-0443',
                'code': 'AP',
                'display': 'Administering Provider'}]},
             'actor': {'reference': 'Practitioner/example-practitioner-id'}}],
           'lotNumber': 'ABC12345',
           'expirationDate': '2026-08-15'}
In [19]: import requests
         import json
```

```
FHIR_BASE = "http://hapi.fhir.org/baseR4"
          patient id = "49233412"
          immunization data = {
            "resourceType": "Immunization",
            "status": "completed",
            "vaccineCode": {
               "coding": [
                 {
                   "system": "http://hl7.org/fhir/sid/cvx",
                   "code": "102",
                   "display": "MMR"
              ],
              "text": "MMR"
            },
            "patient": {
              "reference": f"Patient/{patient_id}"
            "occurrenceDateTime": "2025-08-15T10:00:00Z",
            "lotNumber": "ABC12345"
          response = requests.post(f"{FHIR_BASE}/Immunization", json=immunization_data
          print(f"Status Code: {response.status code}")
          print(response.json())
         Status Code: 201
         {'resourceType': 'Immunization', 'id': '50166507', 'meta': {'versionId': '1'
, 'lastUpdated': '2025-10-14T19:33:59.477+00:00', 'source': '#ipCD4S7Yi0VF7q
         0j'}, 'status': 'completed', 'vaccineCode': {'coding': [{'system': 'http://h
         l7.org/fhir/sid/cvx', 'code': '102', 'display': 'MMR'}], 'text': 'MMR'}, 'pa
         tient': {'reference': 'Patient/49233412'}, 'occurrenceDateTime': '2025-08-15
         T10:00:00Z', 'lotNumber': 'ABC12345'}
In [20]: FHIR_BASE = "http://hapi.fhir.org/baseR4"
          immunization_id = "50166507"
          response = requests.get(f"{FHIR_BASE}/Immunization/{immunization_id}")
          print(response.json())
         {'resourceType': 'Immunization', 'id': '50166507', 'meta': {'versionId': '1'
, 'lastUpdated': '2025-10-14T19:33:59.477+00:00', 'source': '#ipCD4S7Yi0VF7q
         0j'}, 'status': 'completed', 'vaccineCode': {'coding': [{'system': 'http://h
         l7.org/fhir/sid/cvx', 'code': '102', 'display': 'MMR'}], 'text': 'MMR'}, 'pa
         tient': {'reference': 'Patient/49233412'}, 'occurrenceDateTime': '2025-08-15
         T10:00:00Z', 'lotNumber': 'ABC12345'}
```