

Using open source to bring clarity to cancer

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COTA

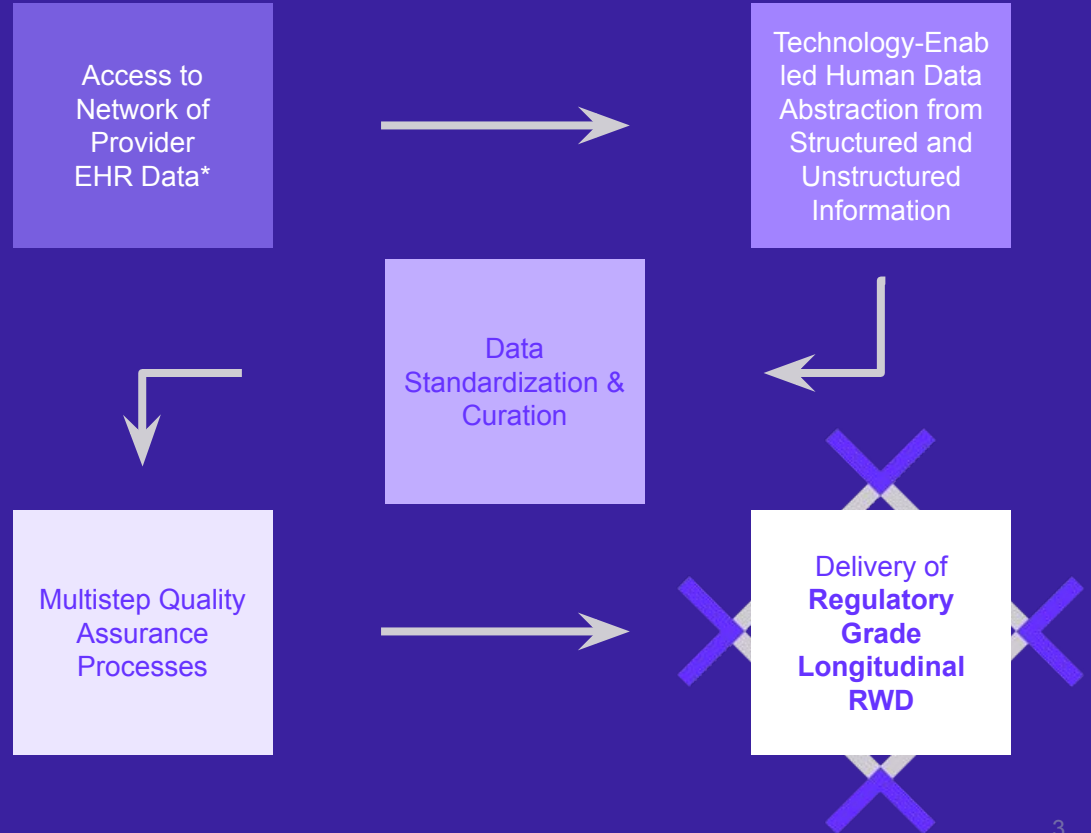
Agenda

- COTA overview and RWA introduction
- How open source has helped us?
- RWA tech stack
- Angular+Plotly.js example
- Cube.js
 - Pre-aggregations example
 - Boolean logic example
 - Asynchronous example

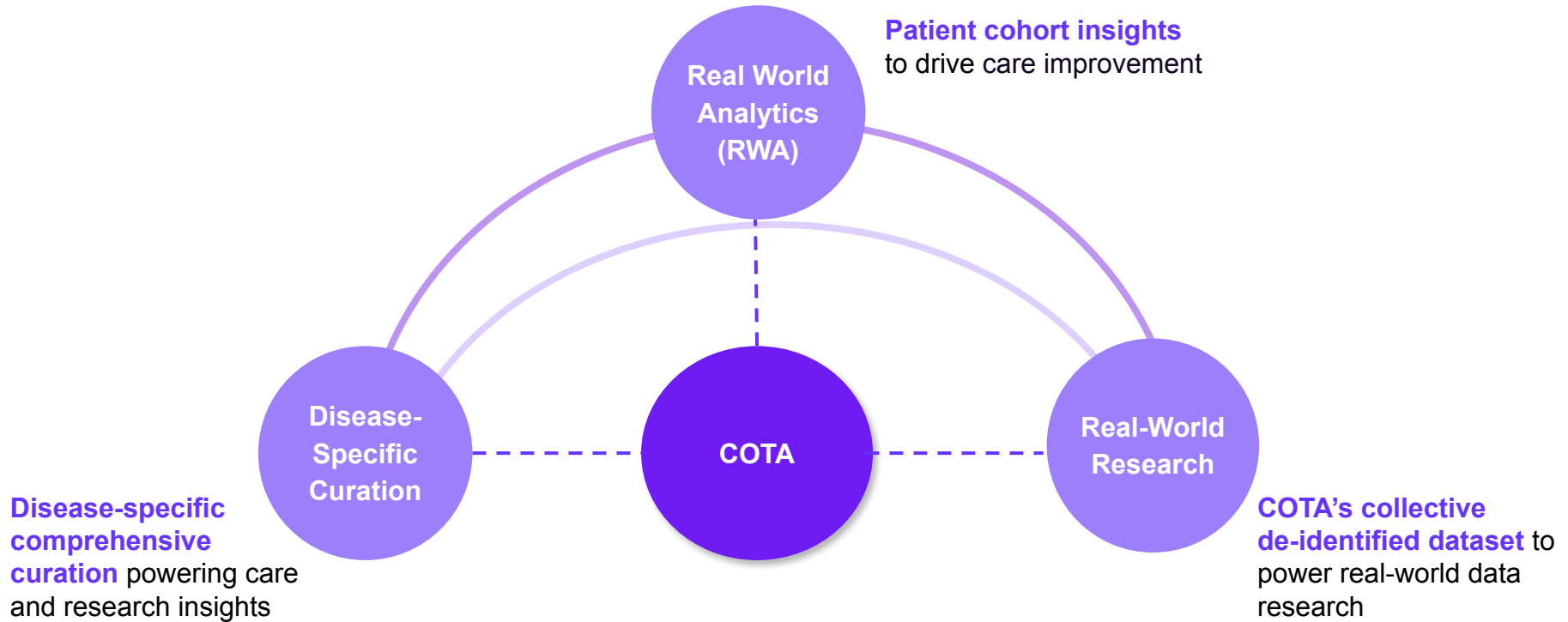
COTA Overview

COTA is a healthcare technology and services company founded in 2011 by doctors, engineers, and data scientists to create clarity from fragmented and often-inaccessible real-world data.

By using our proprietary technology, advanced analytics, and deep expertise to organize complex data, we provide a comprehensive picture of cancer that can be used to advance patient care and research.



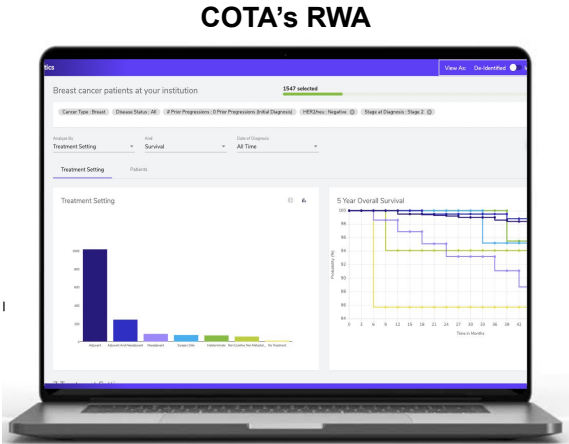
Powered by our platform, COTA offers providers a suite of solutions for not only research, but also to measure and track care delivery and quality



COTA's RWA is a web application that provides the following core tools to answer key questions for hospitals

Provider
Systems

1. Normalization, classification
and enrichment



2. Query and
explore

3. Compare
treatments and
associated
outcomes

4. Track key
operational
metrics and
clinical insights

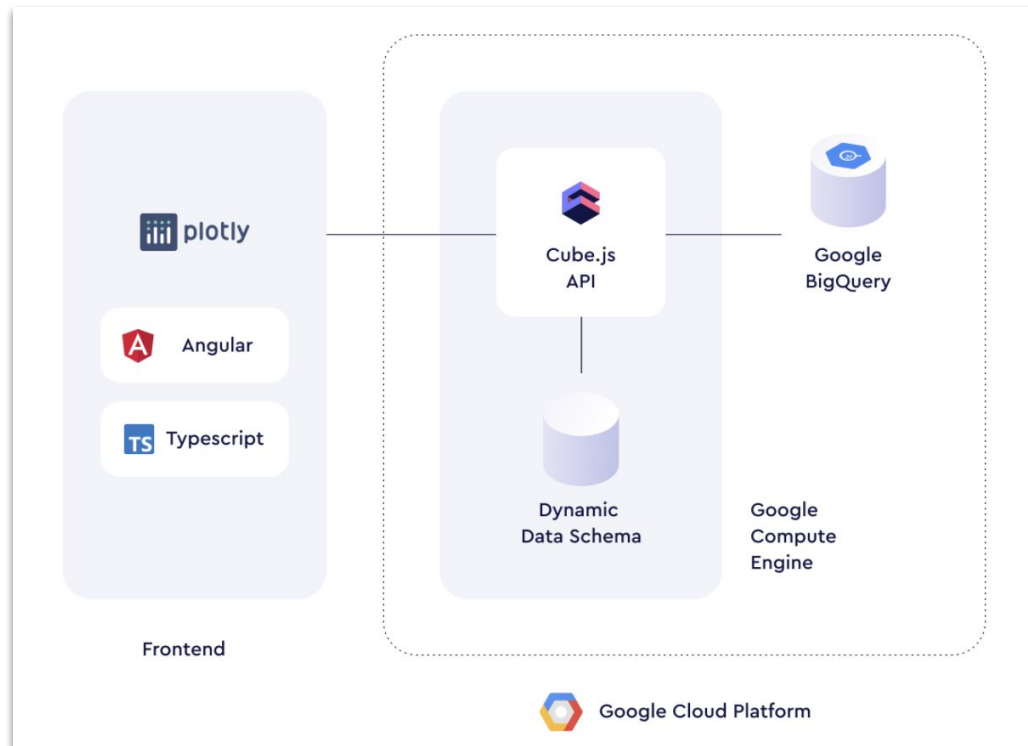
5. Longitudinal
patient timeline
and dashboard

How has open source helped us?

- Moved from using proprietary product to open source, we realised the following advantages :
 - Cost effective
 - Faster development
 - Customization possible
 - No vendor lock-ins, so update or replace if not supporting your cause
 - Better support and quicker iterations

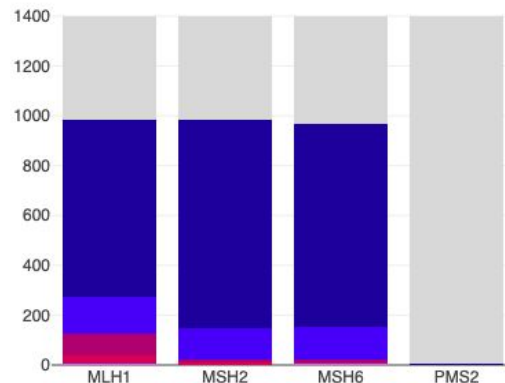
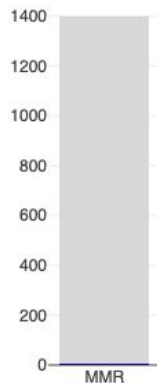
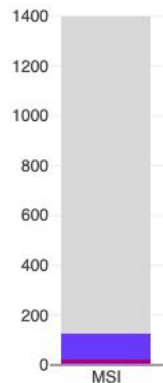
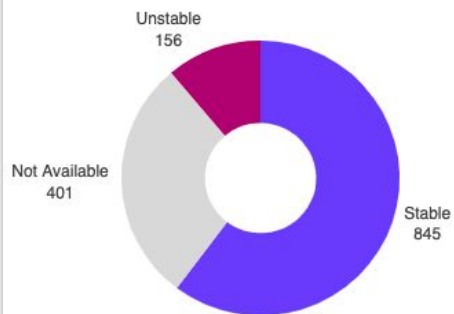
RWA Tech Stack

- Angular
- Plotly
- Cube.js
- Codecept
- Node.js
- TypeScript
- Libraries
 - Moment
 - Lodash
- And many more



Angular / Plotly.js - Subplots

MSI/MMR Status



MSI/MMR Status

- Not Available
- Unstable
- Stable

MSI

- Not Available
- Stable
- High
- Low
- Inconclusive

MMR

- Not Available
- Proficient

Cube.js - Flexibility, support, community

- COTA RWA was seeded along same time as Cube.js was setting up its foot in the OS world. COTA RWA and Cube.js has grown up together!
- Community effect was demonstrated since early stage. With limited capability at start, both the products influenced each-other and benefit from it.
- Contributions were done from product perspective and issues / nice-to-have-features were reported.
- Cube.js community is growing. Support from and within the community is awesome!

Cube.js - Pre-Agg

Data Massaging in Service Layer

```
const calcBMI = (height, weight) => {
  if (!height) {
    return null;
  }
  return (weight/height/height)*703;
}

setBMIBucket = (height, weight) => {
  const bmi = calcBMI(height, weight);
  if (bmi < 18.5) {
    return '< 18.5';
  } else if (bmi >= 18.5 AND bmi < 24.95) {
    return '18.5 - 24.9';
  } else if (bmi >= 24.95 AND bmi < 29.95) {
    return '25 - 29.9';
  } else if (bmi >= 29.95) {
    return '> 30';
  }
  return 'Not Available';
}
```

Pre Agg in Data Layer

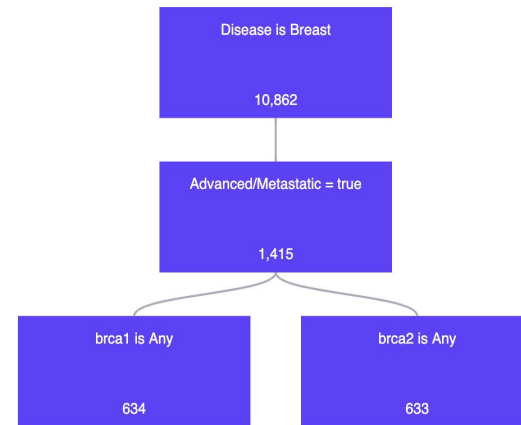
```
WITH
  BMI_CALCULATIONS AS (
    SELECT
      id,
      CASE
        WHEN CAST(height AS FLOAT64) = 0.0 THEN NULL
      ELSE
        CAST(weight AS FLOAT64)/CAST(height AS FLOAT64)/CAST(height AS FLOAT64)*703
      END
      AS bmi
    FROM main_table )
SELECT
  *,
  CASE
    WHEN bmi < 18.5 THEN '< 18.5'
    WHEN bmi >= 18.5 AND bmi < 24.95 THEN '18.5 - 24.9'
    WHEN bmi >= 24.95 AND bmi < 29.95 THEN '25 - 29.9'
    WHEN bmi >= 29.95 THEN '> 30'
  ELSE
    'Not Available'
  END
  AS bmi_bucket
FROM
  main_table main
LEFT JOIN
  BMI_CALCULATIONS calc
ON
  main.id = calc.id
```

Cube.js - Boolean Logic

Metastatic breast cancer patient, should have minimum 1 BRCA (a specific marker) test

▼ And Or + ⊕

Disease	is	Breast	-	
Advanced/Metastatic	=	<input checked="" type="checkbox"/>	-	
▼ <input type="radio"/> And <input checked="" type="radio"/> Or + ⊕ ⊖				
Test Name	is	BRCA1	Any	-
Test Name	is	BRCA2	Any	-



Total Records 640

Cube.js - Async

Static Schema

```
cube(`Test`, {
  sql: `SELECT * FROM Test`,

  sqlAlias: `test`,

  joins: {},

  measures: {},

  dimensions: {
    id: {
      sql: `id`,
      type: `string`,
      primaryKey: true
    },

    testA: {
      sql: `test_a`,
      type: `string`
    },

    testB: {
      sql: `test_b`,
      type: `string`
    },

    testC: {
      sql: `test_c`,
      type: `string`
    }
  }
});
```

Dynamic Schema

```
asyncModule(async () => {
  // columns in Test table : ['id', 'attribute'] // 1 | {test_a: positive, test_b: negative}.....
  const attributeKeys = ['test_a', 'test_b']

  cube(`Test`, {
    sql: `SELECT * FROM Test`,

    measures: {
      count: { sql: `id`, type: `countDistinct` }
    },

    dimensions: {
      id: { sql: `id`, type: `number` },
      attribute: { sql: `attribute`, type: `string` }
    },

    attributeKeys
      .map((key) => ({
        [`${getDimensionName(key)}`]: { // eg: test_a => testA
          sql: `JSON_VALUE(${CUBE}.attribute, '${key}')`,
          type: `string`,
        },
      }))
      .reduce((a, b) => Object.assign(a, b))
  });
});
```

Thank you!