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# Title: Quantifying the Spectrum of Early Motor and Language Milestones in Sex **Chromosome Trisomy** Talia Thompson, PhD<sup>1,2,3</sup>, Samantha Bothwell, M.S.<sup>1,3</sup>, Jennifer Janusz, PsyD<sup>1,2</sup>, Rebecca Wilson, PsyD<sup>1,2</sup>, Susan Howell, MBA, MS, CGC<sup>1,2</sup>, Shanlee Davis, MD, PhD<sup>1,2</sup>, Karli Swenson, PhD, MPH<sup>1,2</sup>, Sydney Martin, MS, OTR<sup>1</sup>, Karen Kowal, PA<sup>4</sup>, Chijioke Ikomi, MD<sup>4</sup>, Maria Despradel<sup>1</sup>, Judith Ross, MD<sup>4</sup>, Nicole Tartaglia, MD<sup>1,2</sup> **Affiliations:** <sup>1</sup>eXtraOrdinarY Kids Clinic & Research Program, Children's Hospital Colorado, Aurora, CO <sup>2</sup>Department of Pediatrics, University of Colorado School of Medicine, Aurora, CO <sup>3</sup>Child Health Biostatistics Core, University of Colorado School of Medicine, Aurora, Colorado. <sup>4</sup>Nemours Children's Hospital, Delaware **ORCID:** Talia Thompson: 0000-0001-6512-9743 Samantha Bothwell: 0000-0002-1616-8137 Jennifer Janusz: 0000-0002-6877-0947 Susan Howell: 0000-0001-7115-2165 Shanlee Davis: 0000-0002-0304-9550 Karli Swenson: 0000-0003-0513-7308 Sydney Martin: 0009-0006-5671-0142 Maria Despradel: 0009-0005-7413-1649 Judith Ross: 0000-0002-8613-2498 Nicole Tartaglia: 0000-0002-8529-6722 Chijioke Ikomi: 0000-0002-0630-2355 **Disclosures:** The authors have no disclosures to report. **Funding:** This study was funded by the eXtraordinarY Babies Study: Natural History of Health and Neurodevelopment in Infants and Young Children with Sex Chromosome Trisomy (NIH NICHD R01HD091251, 3R01HD091251-05S1) **Corresponding Author:** Nicole Tartaglia, MD 13123 East 16<sup>th</sup> Ave, B140 Aurora, CO 80045 Nicole.Tartaglia@ChildrensColorado.org **Author contributions:** Conceptualization - Talia Thompson, Nicole Tartaglia Statistical analysis - Samantha Bothwell

45 - Data collection - Jennifer Janusz, Rebecca Wilson, Susan Howell, Nicole Tartaglia,
 46 Shanlee Davis, Judy Ross, Karen Kowal, Chijioke Ikomi, Talia Thompson

47 48 49 50	<ul> <li>Critical reviewing of the manuscript - Jennifer Janusz, Rebecca Wilson, Susan Howell, Shanlee Davis, Karli Swenson, Sydney Martin, Judith Ross, Maria Despradel</li> <li>Data cleaning - Maria Despradel</li> </ul>
51	Abbreviations:
52	SCT - Sex Chromosome Trisomy
53	CDC - Centers for Disease Control
54	AAP - American Academy of Pediatrics
55	DS - Down syndrome
56	FXS - Fragile X syndrome
57	CNS - Central Nervous System
58	WHO - World Health Organization
59	PRP - Primitive Reflex Profile
60	SDoH - Social Determinants of Health
61	EI – Early Intervention
62	
63	Abstract:
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65	Background and objectives: Sex chromosome trisomy (SCT) is a common chromosomal
66	abnormality associated with increased risks for early developmental delays and
67	neurodevelopmental disorders later in childhood. Our objective was to quantify the spectrum of
68	early developmental milestones in SCT. We hypothesized later milestone achievement in SCT
69	than the general population.
70	
71	Methods: Data were collected as part of the eXtraordinarY Babies Study, a prospective natural
72	history of developmental and health trajectories in a prenatally identified sample of infants with
73	SCT. Parent reported, clinician-validated, early motor and language milestones were collected at
74	2, 6, 12, 18, 24, and 36-months. Age distributions of milestone achievement were compared with
75	normative data.
76	
77	<b>Results</b> : In all SCT conditions, compared with normative data, there was increased variability
78	and a later median age of skill development across multiple gross motor and expressive language
79	milestones. Results also show a significant amount of overlap with the general pediatric
80	population, suggesting that for many children with prenatally identified SCT, early milestones
81	present within, or close to, the expected timeline.
82	
83	<b>Conclusions</b> : As increasing numbers of infants with prenatal SCT diagnoses present at pediatric
84	practices, we provide an evidence-based schedule of milestone achievement in SCT as a tool for
85	pediatricians and families. Detailed data on SCT milestones can support clinical interpretation of
86	milestone achievement. Increased variability and later median age of milestone acquisition in
87	SCT compared to norms support consideration of all infants with SCT as high risk.
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91	Background
92	Sex chromosome trisomy (SCT) (XXY/Klinefelter syndrome, XYY/Jacob syndrome,
93	XXX/Trisomy X) is a common chromosomal abnormality, occurring in 1 of every 500 live
94	births. <sup>1</sup> Prior SCT research, limited by ascertainment bias and small sample sizes, has provided
95	broad descriptions of early development, including profiles of increased risk for delays in gross
96	motor and communication <sup>2,3</sup> and high rates of early intervention. <sup>4</sup> Recent advances in
97	noninvasive prenatal screening <sup>5</sup> have led to increasing rates of prenatally identified SCT and
98	subsequently a growing population of infants with a confirmed SCT diagnosis early in life. As
99	the literature lacks concrete information on the timing of typical milestone achievement in SCT,
100	parents and providers lack clear guidance on what to expect during a child's early years.
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102	Close surveillance of key developmental milestones is a critical part of pediatric care, supporting
103	the promotion of healthy development and the early detection of potential developmental
104	delays. <sup>6</sup> However, common surveillance methods (e.g., CDC milestones <sup>7</sup> checklists) may have
105	less utility for children with genetic conditions and those at-risk for delays such as infants born
106	prematurely. Research has shown that the timing of milestone acquisition differs from the
107	general population in children with Down syndrome (DS), <sup>8</sup> fragile X (FXS), <sup>9</sup> and preterm and
108	very low birthrate infants. <sup>9-11</sup> If this is the case for SCT as well, early developmental care should
109	go beyond surveillance and general screening to include periodic direct developmental
110	assessment. Further, a clear understanding of when children with SCT acquire key
111	developmental milestones is critical for setting reasonable expectations, alerting families to
112	potential concerns, and guiding providers in their referrals for early intervention. This is
113	especially important with the increased frequency of prenatal SCT diagnoses, as pediatricians

114	will be responsible for developmental care in a higher number of infants with SCT presenting to
115	their practices. Therefore, the primary purpose of this study is to fill this gap in the SCT literature
116	with a current, evidence-informed schedule of key early gross motor and language milestone
117	achievement for each of the SCT conditions. These findings will support a more personalized
118	approach to monitoring and care in SCT. Comparisons with previously published normative data
119	to the three SCT conditions will provide critical context and a richer understanding of the SCT
120	phenotypes, and guide recommendations for early developmental care.
121	Methods
122	Data were collected as part of the IRB-approved eXtraordinarY Babies Natural History Study,
123	which leverages recent advances in genetic testing with a prospective investigation of the
124	developmental and health trajectories in a prenatally identified sample of infants with SCT
125	(ClinicalTrials.gov NCT03396562; COMIRB 17-0118; Nemours IRB# 1151006). Inclusion
126	criteria are prenatal identification of SCT (by cfDNA, chorionic villi sampling, and/or
127	amniocentesis) with diagnostic confirmatory karyotype (chorionic villi sampling, amniocentesis,
128	or postnatal), English or Spanish speaking, and child age of 6 weeks to 12 months upon
129	enrollment. Children are excluded from participation if there is a previous diagnosis of a
130	different genetic or metabolic disorder with neurodevelopmental or endocrine involvement,
131	prematurity less than 34 weeks gestational age, a complex congenital malformation not
132	previously associated with SCT, history of significant neonatal complications (i.e.,
133	intraventricular hemorrhage, meningitis, hypoxic-ischemic encephalopathy), or known central
134	nervous system (CNS) malformation identified by neuroimaging. Study visits are conducted
135	regularly at ages 2, 6, 12, 18, 24 months, and then yearly at two sites (Colorado and Delaware)
136	with a combination of in-person and telehealth visits. Visits include comprehensive health and

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137	developmental history, current interventions, physical examination, and a battery of
138	developmental assessments and parent questionnaires. Participants with gestational age <37
139	weeks at birth were excluded from this analysis. Tartaglia et al., (2020) provides additional
140	details on the eXtraordinarY Babies natural history study protocol.
141	
142	Developmental Milestone Measurement
143	Data on the timing of milestones were collected at every study visit as part of a parent completed
144	health and development questionnaire asking parents to report the age (in months) their child
145	achieved key developmental milestones, including eight gross motor skills (rolling front to back,
146	rolling back to front, sitting independently, crawling, cruising, walking, running, jumping) and
147	four expressive communication milestones (cooing, babbling, single words, 2-word phrases).
148	These milestones were chosen because they can be easily observed by parents within a natural
149	setting and delays may predict other areas of known concern in older children with SCT. During
150	the study visit, a physician then reviewed the parent questionnaire responses by interview to
151	confirm ages and parent understanding of the milestone. If there were discrepancies between

152 parent reported skill and the milestone achieved (for example the parent reported the infant was

153 "sitting independently" but physician confirmed the infant was only sitting in a propped

154 position), the physician would adjust the data on the physician data form. The physician data

155 156

157 *Normative data* 

form was used for data analysis.

Each of the twelve developmental milestones collected for the study sample were compared withexisting published norms. We included normative data from studies with published values for the

160	25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> , and 90 <sup>th</sup> percentiles for the milestones of interest from the Denver II Scales, <sup>12</sup> the
161	World Health Organization (WHO) Motor Development Study, <sup>13</sup> and the Primitive Reflex Profile
162	(PRP). <sup>14</sup> As normative data were not available from a single source for all twelve milestones, we
163	used the Denver II whenever possible (sitting, walking, running, jumping, cooing, babbling,
164	single words, 2-word phrases). For milestones that were not included on the Denver II, we used
165	data from the WHO (crawling and cruising) and the PRP (rolling front to back, rolling back to
166	front). As the PRP normative dataset only provided means and standard deviations, percentiles
167	were estimated theoretically under the assumption of a normal distribution.
168	Analysis
169	All analyses were performed in R, version 4.4.0. Descriptive summaries by SCT are presented as
170	median [interquartile range] and N (%). Demographic differences between SCTs were tested
171	using Kruskal-Wallis tests for continuous variables and Fisher's-Exact tests for categorical
172	variables. For each milestone, achieved ages earlier than the normative 2.5 <sup>th</sup> percentile were
173	removed as early outliers. Normative and SCT milestone ages are visualized from their $25^{th}$ –
174	$50^{\text{th}}$ , $50^{\text{th}} - 75^{\text{th}}$ , and $75^{\text{th}} - 90^{\text{th}}$ percentiles. Differences in milestones were analyzed using
175	simulated data based on the normative percentiles, under the assumption of a non-normal
176	distribution, and were tested with Wilcoxon Rank-Sum tests. Differences in milestones were also
177	analyzed between children who had a history of early intervention (EI) therapies and those who
178	did not. Exploring whether there was an overall relationship of milestone achievement with
179	receiving EI therapy was important to ensure therapies were not significantly impacting the
180	distribution of milestones achievement.
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183	Participants include 298 young children with prenatally identified SCT, including 174 with XXY,
184	50 with XYY, and 74 with XXX. All included children had at least one milestone age reported.
185	Table 1 shows sample characteristics. At the time of analysis, the median age of patients included
186	was 4.5 years with the youngest group being XYY children, with a median age of 2.6. The
187	majority of the cohort was white (81.9%) and non-Hispanic/Latinx (83.9%). Included children
188	had participated in the eXtraordinarY Babies study for a median of 3 years, with XXY children
189	having participated the longest (median: 3.5 [IQR: 2.7, 3.8] years) and XYY children having
190	participated for the shortest period of time (median: 1.2 [IQR: 0.4, 3.1] years).
191	
192	Timing of Milestone Achievement in SCT Compared with Normative Datasets.
193	Figure 1 depicts the age (in months) of milestone achievement for each SCT compared with
194	reference norms. Age distributions are characterized by plotting the values for the 25 <sup>th</sup> , 50 <sup>th</sup> , 75 <sup>th</sup> ,
195	and 90 <sup>th</sup> percentiles of each milestone and comparing to normative data. Results indicate that the
196	distributions for all twelve milestones differ (later median milestone achievement; p<0.05) from
197	the normative dataset in at least one SCT group per milestone.
198	
199	Group Differences.
200	Table 2 shows statistical results for group differences in age of milestone achievement between
201	the SCT conditions. Results show statistically significant group differences in cooing (p=0.005);
202	boys with XXY achieved cooing earlier than both girls with XXX ( $p = 0.016$ ) and boys with
203	XYY ( $p = 0.006$ ). Overall group differences exist for crawling ( $p=0.050$ ) and cruising ( $p=0.012$ ).

Boys with XXY achieved crawling (p=0.017) and cruising (p=0.006) at a significantly younger

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205	age than girls with XXX. All other milestone data were statistically similar across trisomy
206	conditions.
207	
208	Comparisons with CDC Milestones
209	Table 3 shows the percent of children by SCT condition who did not achieve milestones by the
210	age listed on the CDC milestones checklists (CDC milestones purport to represent the specific

- 211 health supervision visit age when  $\geq$ 75% or more of children are expected to demonstrate the
- 212 skill).
- 213
- 214 Consideration of Early Intervention Therapies
- 215 Of the 298 children included, 187 (63.8%) had received EI therapy, started either proactively due
- 216 to risk for delays or in response to developmental concerns in one or more developmental
- 217 domains. There were no differences in therapy rates between the SCT conditions. Within our
- 218 cohort, children with history of EI achieved milestones significantly later than children who had
- 219 not (p<0.001 for all milestones). This is likely because those with identified delays were more
- 220 likely to be referred for developmental therapies.
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#### Discussion

This study represents the first report on developmental milestone achievement in prenatally identified SCT and provides a novel milestone chart that can help parents and professionals better quantify and visualize what "increased risk for developmental delay" means in SCT conditions. These cohorts were not referred for any concerns and thus were as close to "population based" as possible. In all SCT conditions, there was a later median age of skill

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228 development across multiple gross motor and expressive language milestones than reported in 229 normative datasets. This includes both early milestones such as cooing and rolling, and later 230 milestones including 2-word phrases, walking, and running. Furthermore, there was more 231 variability in the age range for milestone achievement in our sample compared with reference 232 norms, with the range of acquisition for all milestones extending later in life for children with 233 SCT. These findings support the need to consider infants with SCT as a group at increased risk 234 for delays and deserving of closer developmental monitoring given that age of early motor and 235 language milestones have been shown to predict longitudinal outcomes across all developmental domains in the general population and clinical samples.<sup>15-26</sup> 236 237 These results confirm prior research indicating increased risk for developmental delays in children with SCT,<sup>4,27,28</sup> consistent with findings in other genetic disorders where milestone 238 acquisition is different than population norms.<sup>29</sup> However, unlike other genetic conditions such 239 as DS and FXS,<sup>30,31</sup> our results also show a significant amount of overlap with the general 240 241 pediatric population. Figure 1 shows that, for many children with prenatally identified SCT, early 242 milestones present within, or close to, the expected timeline. While this is reassuring, there are 243 known later risks in SCT for many neurodevelopmental diagnoses including speech-language 244 disorders, learning disabilities, ADHD, executive dysfunction, motor skill deficits, and autism spectrum disorders,<sup>32-44</sup> which all benefit from earlier diagnosis and evidence-based treatments. 245 246 Thus, careful attention to development trajectories is warranted as early interventions may help 247 minimize these morbidities.

248 The variability of the phenotype and overlap with the general population often leads to questions 249 of whether different developmental care pathways and extra developmental testing is needed for 250 *all* infants with SCT. This is a valid concern as a relatively high proportion of individuals with

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SCT conditions have minimal neurodevelopmental differences with positive adult outcomes,<sup>45-47</sup> 251 252 and many go undiagnosed from their clinical presentation. Additional recommendations for 253 developmental monitoring and evaluation may increase family stress, negatively impact parent-254 child relationships, and call unnecessary attention to the genetic differences in their child, as well 255 as increase healthcare utilization and demand on a stressed early intervention system. 256 Prospective longitudinal research is needed to clarify if indeed there are specific early risk 257 factors predictive of poorer outcomes that would warrant stratifying children with SCT into 258 different low versus high-risk developmental care pathways, similar to extensive work done in the congenital heart disease and prematurity populations.<sup>48,49</sup> These pathways, however, were 259 260 developed using evidence from hundreds of studies, which do not currently exist in SCT. Thus, 261 until more prospective data is available, consideration of all infants with SCT as high risk is 262 warranted.

263 Table 3 responds to our interest in whether recently published milestones from the  $CDC^7$  are 264 appropriate for developmental surveillance in infants with SCT. Overall, a relatively small 265 proportion of children in our sample were delayed in milestone achievement according to CDC 266 milestones checklists (Table 3) even though their milestone acquisition was delayed as compared 267 to other metrics (Denver II; WHO). This suggests that relying on the CDC milestone lists for 268 SCT will fail to identify many infants with delayed milestones and is consistent with other published concerns<sup>50-52</sup> about low sensitivity of the ages presented in the CDC milestones. It is 269 270 well recognized that standard developmental screening tools designed for the general population (e.g., ASO, PEDS)<sup>53-55</sup> have lower sensitivity in high-risk groups, which has led to guidelines for 271 developmental follow-up of high-risk neonates with periodic direct assessment.<sup>49,56-58</sup> Similarly, 272

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our findings of the increased risk in SCT support that periodic direct developmental assessment
 should be part of SCT treatment guidelines.<sup>59</sup>

275 By offering detailed information on milestone achievement, we provide a valuable tool for 276 clinicians and families to better interpret a child's early development within the context of their 277 SCT condition, rather than only comparing to general population norms. Further, any significant 278 deviations from SCT norms may alert clinicians to potential risks for comorbid health conditions 279 or an additional genetic difference. While pediatric providers can use this tool as a reference to 280 contextualize a child's milestone achievement, it is not intended to delay referrals for 281 developmental evaluations or early intervention support. Parents may appreciate the more 282 nuanced normative data as they track their child's milestones, noting areas where their child's 283 development aligns with children with similar genetic profiles as well as areas of normative 284 differences. Prior research shows parents of children with delayed milestones may have higher levels of perceived stress<sup>60</sup> or experience guilt that they have done something to cause their 285 child's delays.<sup>61</sup> A clearly defined schedule for the timing of developmental milestones specific 286 287 to each SCT, when used in conjunction with normative milestones expectations, may be more 288 palatable in supporting early developmental care.

Results showing similarities and differences in milestone achievement by karyotype (Table 2) add to the existing literature on genetic disorders by providing more specific data regarding milestone acquisition in each trisomy condition. For most milestones, SCT groups were statistically similar. This aligns with prior research showing similar early developmental and neurocognitive profiles across the SCT conditions.<sup>44,62-64</sup> However, the XXY group did achieve several milestones earlier, including cooing 1 month earlier than both other groups and crawling and cruising 1 month earlier than those with XXX. While this may be an artifact of a larger and

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more variable sample size in XXY, it may also reflect differential effects of the extra X
chromosome in males.<sup>65</sup> Ongoing research with larger sample sizes for XYY and XXX will help
determine if different SCT conditions have clinically relevant differences in developmental
trajectories.

These study results also have practical implications for genetic counseling and are responsive to 300 301 prior research findings showing that parents receiving a prenatal SCT diagnosis desire more 302 accurate and current data on potential neurodevelopmental outcomes specific to each SCT 303 condition. In the context of highly variable phenotypes associated with SCT, genetic counselors 304 strive to provide guidance to parents with a new diagnosis and clarify parental perception of risks for developmental delays.<sup>66</sup> This foundation establishes how parents understand and respond to 305 306 their child's development and behavior, especially as related to the genetic diagnosis. By 307 providing a clearer picture of developmental expectations associated with the diagnosis, genetic 308 counselors can more specifically inform parents about what to expect in their child's first few 309 years of life, as well as promote awareness, empowerment, and a proactive approach to early intervention processes to facilitate early developmental care.<sup>67</sup> 310

311 Despite the insights gained, limitations are important to consider. First, smaller sample sizes for 312 the XXX and XYY karyotypes limit the generalizability compared with the XXY sample. 313 Additionally, there are known limitations in normative data for milestone acquisition, including unclear and inconsistent definitions of milestones,<sup>68</sup> ambiguity around what constitutes 314 achievement of the milestone (partial vs. complete),<sup>69</sup> and differences in the raters used to 315 determine milestone achievement for normative datasets (parents vs. clinicians).<sup>68,70,71</sup> Further, 316 normative datasets rarely account for potential sex differences,<sup>72,73</sup> racial and sociocultural 317 differences, <sup>50,74,75</sup> and known variability related to social determinants of health<sup>69</sup> (SDoH), which 318

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319 further adds a degree of uncertainty to our findings. Our sample was disproportionately white, 320 non-Hispanic with high socioeconomic status, and future studies should aim to include more 321 representative samples. Parental recall bias is another commonly recognized challenge when evaluating parent-reported milestones,<sup>76,77</sup> however minimized in this study with frequent visits 322 323 at 2, 6, 12, 18, 24, and 36 months of age with pediatric physicians interviewing and verifying 324 milestone achievements. Importantly, while there are many benefits to an ongoing natural history 325 study, our study design is limited in that at the time of publication, not all participants in the 326 sample had yet achieved all 12 milestones measured and therefore sample sizes were different 327 for each SCT condition at each milestone. Additionally, while we explored the effect of early 328 intervention in our analysis, the act of participating in a natural history study itself may impact 329 developmental course. Families in the study have self-selected to participate in regularly 330 scheduled developmental evaluations with expert SCT clinicians and to monitor developmental 331 milestones in between visits. While a prenatally identified sample of nearly 300 infants with SCT 332 provides a less biased dataset than prior studies, it may still not fully represent the broad 333 spectrum of outcomes in SCT. Future results based on direct assessments through the 334 eXtraordinarY Babies study can address these limitations and further refine our understanding of 335 developmental trajectories and risk groups in this population.

In conclusion, developmental milestone achievement in SCT conditions is delayed compared to the general population, however only in a subset of infants with SCT. As increasing numbers of infants with prenatal SCT diagnoses present at pediatric practices, we provide an evidence-based schedule of milestone achievement in SCT as a tool for families, pediatricians, genetic counselors, and early intervention teams. Utilization of such a tool can support shared clinical decision-making between parents and providers, promoting timely referrals and identifying

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342	patterns inconsistent with SCT. However, given the paucity of prospective research identifying
343	specific risk factors for later negative outcomes, recommended care for SCT conditions should
344	follow practices of other high-risk conditions - with more responsive attention to developmental
345	concerns, recognition that standard surveillance and screening tools have lower sensitivities in
346	high-risk populations, and referrals for periodic direct developmental assessments. While more
347	rigorous research will help identify evidence for timing of direct assessments and highest-risk
348	groups, general publications support assessments at 6-12 months, 18-24 months, and 36 months
349	of age. <sup>78-80</sup>

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## 351 Table 1. Cohort Demographics

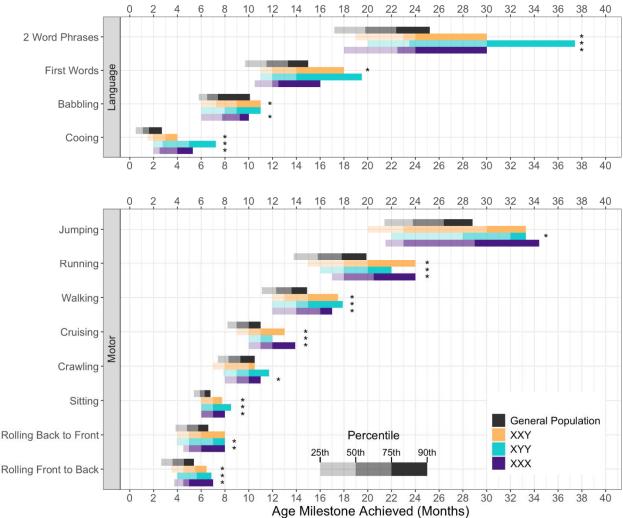
	Overall (N=298)	XXY (N=174)	XYY (N=50)	XXX (N=74)	P-Value
Age (Years; As of 7/9/2024)					
Median [IQR]	4.5 [2.9, 5.7]	4.9 [3.8, 6.1]	2.6 [1.4, 5.3]	3.5 [2, 5.1]	< 0.001*
Years in Study					
Median [IQR]	3 [1.4, 3.8]	3.5 [2.7, 3.8]	1.2 [0.4, 3.1]	2.5 [0.9, 3.5]	< 0.001*
Race					
White	244 (81.9%)	138 (79.3%)	41 (82.0%)	65 (87.8%)	0.119
Native Hawaiian or Other Pacific Islander	1 (0.3%)	1 (0.6%)	0 (0%)	0 (0%)	
African American or Black	17 (5.7%)	13 (7.5%)	4 (8.0%)	0 (0%)	
Asian	24 (8.1%)	14 (8.0%)	2 (4.0%)	8 (10.8%)	
Native American or Alaska Native	3 (1.0%)	2 (1.1%)	1 (2.0%)	0 (0%)	
Other	6 (2.0%)	5 (2.9%)	1 (2.0%)	0 (0%)	
Missing	3 (1.0%)	1 (0.6%)	1 (2.0%)	1 (1.4%)	
Ethnicity					
Hispanic/Latinx	45 (15.1%)	28 (16.1%)	6 (12.0%)	11 (14.9%)	0.859
Non-Hispanic/Latinx	250 (83.9%)	145 (83.3%)	43 (86.0%)	62 (83.8%)	
Missing	3 (1.0%)	1 (0.6%)	1 (2.0%)	1 (1.4%)	
Hollingshead Index					
Median [IQR]	54.5 [47.9, 59.5]	54 [47, 59.5]	54.5 [46.1, 59.2]	55.5 [50.5, 59.5]	0.619
Missing	10 (3.4%)	1 (0.6%)	4 (8.0%)	5 (6.8%)	
Annual Family Income*					
\$50,000 or less	17 (5.7%)	12 (6.9%)	3 (6.0%)	2 (2.7%)	0.437
\$50,000 - \$100,000	67 (22.5%)	38 (21.8%)	15 (30.0%)	14 (18.9%)	
\$100,000 - \$250,000	154 (51.7%)	91 (52.3%)	20 (40.0%)	43 (58.1%)	
> \$250,000	54 (18.1%)	30 (17.2%)	11 (22.0%)	13 (17.6%)	
Missing	6 (2.0%)	3 (1.7%)	1 (2.0%)	2 (2.7%)	

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Significance level = 0.05. Overall differences tested using Kruskal-Wallis tests for continuous variables and Fisher's Exact/Chi-Squared Tests for categorical variables. \*Family income data reported were collected at initial eXtraordinarY Babies study visit.

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### 356 Figure 1. Achievement of Language and Motor milestones in Sex Chromosome Trisomy



### 357 compared to the general population

Age Milestone Achieved (Months) \*: Trisomy is delayed compared to simulated data based on general population percentiles, under the assumption of a non-normal

distribution. Significance level = 0.05. Differences tested with Wilcoxon Rank-Sum Tests.

361 General Population estimates are based on Denver II for Jumping, Running, Walking, Sitting, 2 Word Phrases, First Words,

362 Babbling, and Cooing; WHO for Cruising and Crawling; and PRP for Rolling Back to Front and Rolling Front to Back.<sup>12-14</sup>

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	XXY (N=174)	XYY (N=50)	XXX (N=74)	Overall Kruskal-Wallis P-Value
Language				
Cooing	N = 165	<i>N</i> = 46	<i>N</i> = <i>6</i> 8	
Median [IQR]	2 [1.5, 3]	2.8 [2, 5]	2.5 [2, 4]	$0.005^{*}$ <sup>1, 2</sup>
Babbling	<i>N</i> = <i>158</i>	N = 44	<i>N</i> = <i>61</i>	
Median [IQR]	7 [6, 9]	7.2 [5.9, 9]	7 [5, 9]	0.867
First Words	N = 152	N = 36	<i>N</i> = 55	
Median [IQR]	12 [11, 14]	12 [11, 14]	12 [10.5, 12.5]	0.557
2 Word Phrases	N = 135	<i>N</i> = 23	N = 42	
Median [IQR]	23 [18, 24]	23 [19, 30]	22.5 [18, 24]	0.637
Motor				
Rolling Front to Back	N = 162	N = 44	<i>N</i> = <i>64</i>	
Median [IQR]	4.5 [3.5, 5.5]	4 [4, 5.6]	4.5 [3.8, 5]	0.891
Rolling Back to Front	<i>N</i> = <i>162</i>	<i>N</i> = <i>43</i>	<i>N</i> = <i>64</i>	
Median [IQR]	5 [4, 6]	5 [4, 7]	5 [4.4, 6]	0.193
Sitting	<i>N</i> = <i>164</i>	<i>N</i> = <i>45</i>	<i>N</i> = <i>60</i>	
Median [IQR]	6 [5.4, 7]	6 [6, 7]	6 [5.5, 7]	0.354
Crawling	<i>N</i> = <i>162</i>	N = 44	<i>N</i> = <i>6</i> 2	
Median [IQR]	8 [7, 9.9]	9 [7.9, 10]	9 [8, 10]	$0.050^{*2}$
Cruising	N = 157	N = 41	<i>N</i> = <i>6</i> 2	
Median [IQR]	10 [9, 11]	11 [10, 12]	11 [10, 12]	$0.012^{*2}$
Walking	N = 156	<i>N</i> = <i>32</i>	<i>N</i> = <i>53</i>	
Median [IQR]	13 [12, 15]	14 [12, 15]	14 [12, 16]	0.239
Running	N = 145	N = 27	<i>N</i> = <i>51</i>	
Median [IQR]	18 [15, 20]	18 [16, 20]	18 [17, 20.5]	0.177
Jumping	<i>N</i> = <i>131</i>	<i>N</i> = 22	<i>N</i> = 43	
Median [IQR]	23 [20, 30]	26.2 [22, 31.5]	23 [21.5, 29]	0.374

#### **Table 2** Age in Months of Milestone Achievement by $SCT^{1}$

368 369 \* Significant difference in at least one trisomy; alpha = 0.05

<sup>1</sup> Pairwise test XXY  $\neq$  XYY; Bonferroni adjusted alpha = 0.017 <sup>2</sup> Pairwise test XXY  $\neq$  XXX; Bonferroni adjusted alpha = 0.017

Milestone	Age	XXY (Total N = 174)	XYY (Total N = 50)	XXX (Total N = 74),
Language		-		
Cooing	4 months	16/165 (9.7%)	13/46 (28.3%)	11/68 (16.2%)
Babbling	9 months	35/158 (22.12%)	9/44 (20.5%)	13/61 (21.3%)
First Words	15 months	32/1582 (21.1%)	8/36 (22.2%)	10/55 (18.2%)
2 Word Phrases	24 months	33/135 (24.4%)	9/33 (39.1%)	10/42 (23.8%)
Motor				
Rolling Front to Back	6 months	17/162 (10.5%)	7/44 (15.9%)	9/64 (14.1%)
Sitting Independently	9 months	3/164 (1.8%)	1/45 (2.2%)	2/60 (3.3%)
Cruising	12 months	22/157 (14%)	3/41 (7.3%)	13/62 (21%)
Walking	15 months	33/156 (21.2%)	7/32 (21.9%)	19/53 (35.9%)
Running	24 months	12/145 (8.3%)	1/27 (3.7%)	4/51 (7.8%)
Jumping	30 months	22/131 (16.8%)	6/22 (27.3%)	7/43 (16.3%)

# 374 Table 3. Frequencies of children with SCT delayed in milestones according to ages set by CDC375 Milestones checklists.

N (%; p-value): Number (%) of children in each trisomy that achieved the milestone later than the ages listed on CDC milestone checklists. Total sample sizes differ for each milestone and trisomy.

378 CDC cut points were not available for rolling back to front and crawling

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